## SEQUENCE LISTING

Fuqua, Suzanne Allred, D. Hopp, Torsten A. O'Connell, Peter

<120> Methods and Composition in Breast Cancer Diagnosis and Therapeutics

<130> P02102US2 <140> US 10/052,092 <141> 2002-01-18 <150> US 60/262,990 <151> 2001-01-19 <150> US 60/304,018 <151> 2001-07-09 <160> 49 <170> PatentIn version 3.1 <210> 1 <211> 6450

<400> 1

DNA <213> Human

<212>

gagttgtgcc tggagtgatg tttaagccaa tgtcagggca aggcaacagt ccctggccgt 60 cctccagcac ctttgtaatg catatgagct cgggagacca gtacttaaag ttggaggccc 120 gggageceag gagetggegg agggegtteg teetgggage tgeaettget eegtegggte 180 gccggcttca ccggaccgca ggctcccggg gcagggccgg ggccagagct cgcgtgtcgg 240 cgggacatgc gctgcgtcgc ctctaacctc gggctgtgct ctttttccag gtggcccgcc 300 ggtttetgag cettetgeee tgeggggaea eggtetgeae eetgeeegeg geeaeggaee 360 atgaccatga ccctccacac caaagcatct gggatggccc tactgcatca gatccaaggg 420 aacgagetgg ageeeetgaa eegteegeag eteaagatee eeetggageg geeeetggge 480 gaggtgtacc tggacagcag caagcccgcc gtgtacaact accccgaggg cgccgcctac 540 gagttcaacg ccgcggccgc cgccaacgcg caggtctacg gtcagaccgg cctcccctac 600 ggccccgggt ctgaggctgc ggcgttcggc tccaacggcc tggggggttt ccccccactc 660 aacagegtgt eteegageee getgatgeta etgeaceege egeegeaget gtegeettte 720 ctgcagcccc acggccagca ggtgccctac tacctggaga acgagcccag cggctacacg 780 gtgcgcgagg ccggcccgcc ggcattctac aggccaaatt cagataatcg acgccagggt 840 ggcagagaaa gattggccag taccaatgac aagggaagta tggctatgga atctgccaag 900

gagactcgct actgtgcagt gtgcaatgac tatgcttcag gctaccatta tggagtctgg 960 tcctgtgagg gctgcaaggc cttcttcaag agaagtattc aaggacataa cgactatatg 1020 tgtccagcca ccaaccagtg caccattgat aaaaacagga ggaagagctg ccaggcctgc 1080 cggctccgca aatgctacga agtgggaatg atgaaaggtg ggatacgaaa agaccgaaga 1140 ggagggagaa tgttgaaaca caagcgccag agagatgatg gggagggcag gggtgaagtg 1200 gggtctgctg gagacatgag agctgccaac ctttggccaa gcccgctcat gatcaaacgc 1260 tctaagaaga acagcctggc cttgtccctg acggccgacc agatggtcag tgccttgttg 1320 gatgetgage eccecataet etatteegag tatgateeta ecagaceett eagtgaaget 1380 tcgatgatgg gcttactgac caacctggca gacagggagc tggttcacat gatcaactgg 1440 gcgaagaggg tgccaggctt tgtggatttg accctccatg atcaggtcca ccttctagaa 1500 tgtgcctggc tagagatcct gatgattggt ctcgtctggc gctccatgga gcacccagtg 1560 aagctactgt ttgctcctaa cttgctcttg gacaggaacc agggaaaatg tgtagagggc 1620 atggtggaga tettegaeat getgetgget acateatete ggtteegeat gatgaatetg 1680 cagggagagg agtttgtgtg cctcaaatct attattttgc ttaattctgg agtgtacaca 1740 tttctgtcca gcaccctgaa gtctctggaa gagaaggacc atatccaccg agtcctggac 1800 aagatcacag acactttgat ccacctgatg gccaaggcag gcctgaccct gcagcagcag 1860 caccagegge tggeecaget ecteeteate eteteceaca teaggeacat gagtaacaaa 1920 ggcatggagc atctgtacag catgaagtgc aagaacgtgg tgcccctcta tgacctgctg 1980 ctggagatgc tggacgccca ccgcctacat gcgcccacta gccgtggagg ggcatccgtg 2040 gaggagacgg accaaagcca cttggccact gcgggctcta cttcatcgca ttccttgcaa 2100 aagtattaca tcacggggga ggcagagggt ttccctgcca cagtctgaga gctccctggc 2160 teccaeaegg tteagataat eeetgetgea ttttaeeete ateatgeaee aetttageea 2220 aattctgtct cctgcataca ctccggcatg catccaacac caatggcttt ctagatgagt 2280 ggccattcat ttgcttgctc agttcttagt ggcacatctt ctgtcttctg ttgggaacag 2340 ccaaagggat tccaaggcta aatctttgta acagctctct ttcccccttg ctatgttact 2400 aagegtgagg atteeegtag etetteaeag etgaaeteag tetatgggtt ggggeteaga 2460 taactctgtg catttaagct acttgtagag acccaggcct ggagagtaga cattttgcct 2520 ctgataagca ctttttaaat ggctctaaga ataagccaca gcaaagaatt taaagtggct 2580 cctttaattg gtgacttgga gaaagctagg tcaagggttt attatagcac cctcttgtat 2640 teetatggea atgeateett ttatgaaagt ggtaeaeett aaagetttta tatgaetgta 2700 gcagagtatc tggtgattgt caattcactt ccccctatag gaatacaagg ggccacacag 2760

25136771 -2-

ggaaggcaga teceetagtt ggeeaagaet tattttaaet tgatacaetg cagatteaga 2820 gtgtcctgaa gctctgcctc tggctttccg gtcatgggtt ccagttaatt catgcctccc 2880 atggacctat ggagagcaac aagttgatct tagttaagtc tccctatatg agggataagt 2940 tcctgatttt tgttttatt tttgtgttac aaaagaaagc cctccctccc tgaacttgca 3000 gtaaggtcag cttcaggacc tgttccagtg ggcactgtac ttggatcttc ccggcgtgtg 3060 tgtgccttac acaggggtga actgttcact gtggtgatgc atgatgaggg taaatggtag 3120 ttgaaaggag caggggccct ggtgttgcat ttagccctgg ggcatggagc tgaacagtac 3180 ttgtgcagga ttgttgtggc tactagagaa caagagggaa agtagggcag aaactggata 3240 cagttetgag cacagecaga ettgeteagg tggeeetgea caggetgeag etacetagga 3300 acatteettg cagaceeege attgeetttg ggggtgeeet gggateeetg gggtagteea 3360 gctcttattc atttcccagc gtggccctgg ttggaagaag cagctgtcaa gttgtagaca 3420 gctgtgttcc tacaattggc ccagcaccct ggggcacggg agaagggtgg ggaccgttgc 3480 tgtcactact caggctgact ggggcctggt cagattacgt atgcccttgg tggtttagag 3540 ataatccaaa atcagggttt ggtttgggga agaaaatcct cccccttcct ccccgcccc 3600 gttccctacc gcctccactc ctgccagctc atttccttca atttcctttg acctataggc 3660 taaaaaagaa aggctcattc cagccacagg gcagccttcc ctgggccttt gcttctctag 3720 cacaattatg ggttacttcc tttttcttaa caaaaaagaa tgtttgattt cctctgggtg 3780 accttattgt ctgtaattga aaccctattg agaggtgatg tctgtgttag ccaatgaccc 3840 aggtagetge tegggettet ettggtatgt ettgtttgga aaagtggatt teatteattt 3900 3960 aaaaagtttt tatgtgcact taaatttggg gacaatttta tgtatctgtg ttaaggatat 4020 gcttaagaac ataattettt tgttgetgtt tgtttaagaa geacettagt ttgtttaaga 4080 agcaccttat atagtataat atatatttt ttgaaattac attgcttgtt tatcagacaa 4140 ttgaatgtag taattctgtt ctggatttaa tttgactggg ttaacatgca aaaaccaagg 4200 aaaaatattt agtttttttt tttttttttg tatacttttc aagctacctt gtcatgtata 4260 cagtcattta tgcctaaagc ctggtgatta ttcatttaaa tgaagatcac atttcatatc 4320 aacttttgta tccacagtag acaaaatagc actaatccag atgcctattg ttggatattg 4380 aatgacagac aatcttatgt agcaaagatt atgcctgaaa aggaaaatta ttcagggcag 4440 ctaattttgc ttttaccaaa atatcagtag taatattttt ggacagtagc taatgggtca 4500 gtgggttctt tttaatgttt atacttagat tttcttttaa aaaaattaaa ataaaacaaa 4560

25136771 - 3 -

aaaaatttct aggactagac gatgtaatac cagctaaagc caaacaatta tacag	tggaa 4620
ggttttacat tattcatcca atgtgtttct attcatgtta agatactact acatt	tgaag 4680
tgggcagaga acatcagatg attgaaatgt tcgcccaggg gtctccagca acttt	ggaaa 4740
tctctttgta tttttacttg aagtgccact aatggacagc agatattttc tggct	gatgt 4800
tggtattggg tgtaggaaca tgatttaaaa aaaaaactct tgcctctgct ttccc	ccact 4860
ctgaggcaag ttaaaatgta aaagatgtga tttatctggg gggctcaggt atggt	gggga 4920
agtggattca ggaatctggg gaatggcaaa tatattaaga agagtattga aagta	tttgg 4980
aggaaaatgg ttaattotgg gtgtgcacca aggttcagta gagtccactt ctgcco	ctgga 5040
gaccacaaat caactagete catttacage catttetaaa atggcagett cagtte	ctaga 5100
gaagaaagaa caacatcagc agtaaagtcc atggaatagc tagtggtctg tgtttc	ctttt 5160
cgccattgcc tagcttgccg taatgattct ataatgccat catgcagcaa ttatga	agagg 5220
ctaggtcatc caaagagaag accctatcaa tgtaggttgc aaaatctaac ccctaa	iggaa 5280
gtgcagtctt tgatttgatt tccctagtaa ccttgcagat atgtttaacc aagcca	ıtagc 5340
ccatgccttt tgagggctga acaaataagg gacttactga taatttactt ttgatc	acat 5400
taaggtgttc tcaccttgaa atcttataca ctgaaatggc cattgattta ggccac	tggc 5460
ttagagtact ccttcccctg catgacactg attacaaata ctttcctatt catact	ttcc 5520
aattatgaga tggactgtgg gtactgggag tgatcactaa caccatagta atgtct	aata 5580
ttcacaggca gatctgcttg gggaagctag ttatgtgaaa ggcaaataaa gtcata	cagt 5640
agctcaaaag gcaaccataa ttctctttgg tgcaagtctt gggagcgtga tctaga	ttac 5700
actgcaccat tcccaagtta atcccctgaa aacttactct caactggagc aaatga	actt 5760
tggtcccaaa tatccatctt ttcagtagcg ttaattatgc tctgtttcca actgca	tttc 5820
ctttccaatt gaattaaagt gtggcctcgt ttttagtcat ttaaaattgt tttcta	agta 5880
attgctgcct ctattatggc acttcaattt tgcactgtct tttgagattc aagaaa	aatt 5940
tctattcatt tttttgcatc caattgtgcc tgaactttta aaatatgtaa atgctg	ccat 6000
gttccaaacc catcgtcagt gtgtgtgttt agagctgtgc accctagaaa caacata	actt 6060
gtcccatgag caggtgcctg agacacagac ccctttgcat tcacagagag gtcattg	ggtt 6120
atagagactt gaattaataa gtgacattat gccagtttct gttctctcac aggtgat	caaa 6180
caatgetttt tgtgeactae ataetettea gtgtagaget ettgttttat gggaaaa	aggc 6240
tcaaatgcca aattgtgttt gatggattaa tatgcccttt tgccgatgca tactatt	act 6300
gatgtgactc ggttttgtcg cagctttgct ttgtttaatg aaacacactt gtaaacc	ctct 6360
tttgcacttt gaaaaagaat ccagcgggat gctcgagcac ctgtaaacaa ttttctc	caac 6420

ctatttgatg ttcaaataaa gaattaaact	6450
<210> 2 <211> 1644 <212> DNA <213> Artificial Sequence	
<220> <223> Synthetic Estrogen Receptor L	igand Binding Domain Fusion
<400> 2	
atggtgagca agggcgagga gctgttcacc ggg	
ggcgacgtaa acggccacag gttcagcgtg tcc	
ggcaagctga ccctgaagtt catctgcacc acc	
ctcgtgacca ccctgacctg gggcgtgcag tgc	
cagcacgact tettcaagte egecatgeee gaa	ggctacg tccaggagcg caccatcttc 300
ttcaaggacg acggcaacta caagacccgc gcc	gaggtga agttcgaggg cgacaccctg 360
gtgaaccgca tcgagctgaa gggcatcgac ttc	aaggagg acggcaacat cctggggcac 420
aagctggagt acaactacat cagccacaac gtc	tatatca ccgccgacaa gcagaagaac 480
ggcatcaagg cccacttcaa gatccgccac aaca	atcgagg acggcagcgt gcagctcgcc 540
gaccactacc agcagaacac ccccatcggc gacq	ggccccg tgctgctgcc cgacaaccac 600
tacctgagca cccagtccgc cctgagcaaa gaco	cccaacg agaagcgcga tcacatggtc 660
ctgctggagt tcgtgaccgc cgcccgcatg cccc	
gccaaccttt ggccaagccc gctcatgatc aaac	
teeetgaegg eegaeeagat ggteagtgee ttgt	
teegagtatg ateetaceag accetteagt gaag	
ctggcagaca gggagctggt tcacatgatc aact	
gatttgaccc tccatgatca ggtccacctt ctag	
attggtctcg tctggcgctc catggagcac ccag	
ctcttggaca ggaaccaggg aaaatgtgta gagg	
ctggctacat catctcggtt ccgcatgatg aatc	
aaatctatta ttttgcttaa ttctggagtg taca	catttc tgtccagcac cctgaagtct 1260
ctggaagaga aggaccatat ccaccgagtc ctgg	acaaga tcacagacac tttgatccac 1320
tgatggcca aggcaggcct gaccctgcag cagc	agcacc agcggctggc ccagctcctc 1380
stcatectet eccaeateag geacatgagt aaca	aaggca tggagcatct gtacagcatg 1440

25136771 - 5 -

augegeaaga	acgugguguu	ccccacgac	ctgctgctgg	agacgetgga	cgcccaccgc	1500
ctacatgcgc	ccactagccg	tggaggggca	tccgtggagg	agacggacca	aagccacttg	1560
gccactgcgg	gctctacttc	atcgcattcc	ttgcaaaagt	attacatcac	gggggaggca	1620
gagggtttcc	ctgccacagt	ctga				1644
<210> 3 <211> 123 <212> DNA <213> Hum						
<400> 3	aatcagtttg	tataaactca	ccaaagggta	atattat	<b></b>	60
						60
	tttgtcagac					120
	catgttatgg					180
	ttctcttgga					240
aagtatttct	tcaaataaaa	tgaaagctgg	ttagctttga	aaattttttg	tctaaaagtt	300
tacacgggaa	aaaaattaac	taatttttt	tttccacctg	tgttttcagg	gatacgaaaa	360
gaccgaagag	gagggagaat	gttgaaacac	aagcgccaga	gagatgatgg	ggagggcagg	420
ggtgaagtgg	ggtctgctgg	agacatgaga	gctgccaacc	tttggccaag	cccgctcatg	480
atcaaacgct	ctaagaagaa	cagcctggcc	ttgtccctga	cggccgacca	gatggtcagt	540
gccttgttgg	atgctgagcc	ccccatactc	tattccgagt	atgatcctac	cagacccttc	600
agtgaagctt	cgatgatggg	cttactgacc	aacctggcag	acagggagct	ggttcacatg	660
atcaactggg	cgaagagggt	gccaggtaag	aatgcgaagc	gcagctttta	agagtcaata	720
gcttttcaag	aacttgttgt	gatgtcatgg	gagaaatagt	gggggaaaaa	gaagcaataa	780
catgttatgt	aattggtttc	aaggttacag	gagatgtgtt	cattttcagt	atcaatacac	840
tgtaattttc	caggagatta	ggaaataata	tttttaaatc	agaatctaga	agactgaaat	900
tcttaaattg	acataattta	tttttaaccc	atctcattta	ccaaaaagat	ttagggtgga	960
cactacatgg	taaaactatt	taatagtgta	tgttcacagt	agcagaaact	tttaacacta	1020
aatgaactac	aaaagtttgt	aatattaatg	acctttgttg	aaaacatctc	aattattaat	1080
caaacgattt	tatcttaaaa	agatttttaa	gattcggtgt	ggtggctcgt	gcctgtaatc	1140
ctagcacttt	ttggggctga	ggtgggagga	ttgcttgagc	ccaagagctt	gaggaaagag	1200
aggacagcga	ttctcgtacg	aacggttacg	attctga			1237

25136771 - 6 -

<sup>&</sup>lt;210> 4 <211> 392 <212> DNA

<213> Human

<400> 4 tcgaagagga gggagaatg	t tgaagcacaa	gcgccagaga	gatgatgggg	agggcagggg	60
tgaagtgggg tctgctgga	g acatgagagc	tgccaacctt	tggccaagcc	cgctcatgat	120
caaacgctct aagaagaac	a gcctggcctt	gtccctgacg	gccgaccaga	tggtcagtgc	180
cttgttggat gctgagccc	c ccatactcta	ttccgagtat	gatcctacca	gacccttcag	240
tgaagcttcg atgatgggc	t tactgaccaa	cctggcagac	agggagctgg	ttcacatgat	300
caactgggcg aagagggtg	c caggaaccag	ggaaaatgtg	tagagggcat	ggtggagatc	360
ttcgacatgc tgctggcta	c atcatctcgg	tt			392
<210> 5 <211> 2106 <212> DNA <213> Human					
<400> 5 atgaccatga ccctccaca	c caaagcatct	gggatggccc	tactgcatca	gatccaaggg	60
aacgagctgg agcccctga	a ccgtccgcag	ctcaagatcc	ccctggagcg	gcccctgggc	120
gaggtgtacc tggacagca	g caagcccgcc	gtgtacaact	accccgaggg	cgccgcctac	180
gagttcaacg ccgcggccg	c cgccaacgcg	caggtctacg	gtcagaccgg	cctcccctac	240
ggccccgggt ctgaggctg	c ggcgttcggc	tccaacggcc	tggggggttt	cccccactc	300
aacagcgtgt ctccgagcc	c gctgatgcta	ctgcacccgc	cgccgcagct	gtcgcctttc	360
ctgcagcccc acggccagc	a ggtgccctac	tacctggaga	acgagcccag	cggctacacg	420
gtgcgcgagg ccggcccgc	c ggcattctac	aggccaaatt	cagataatcg	acgccagggt	480
ggcagagaaa gattggcca	g taccaatgac	aagggaagta	tggctatgga	atctgccaag	540
gagactcgct actgtgcag	t gtgcaatgac	tatgcttcag	gctaccatta	tggagtctgg	600
tcctgtgagg gctgcaagg	c cttcttcaag	agaagtattc	aaggacataa	cgactatatg	660
tgtccagcca ccaaccagt	g caccattgat	aaaaacagga	ggaagagctg	ccaggcctgc	720
cggctccgca aatgctacg	a agtgggaatg	atgaaaggtg	ggatacgaaa	agaccgaaga	780
ggagggagaa tgttgaaac	a caagcgccag	agagatgatg	gggagggcag	gggtgaagtg	840
gggtctgctg gagacatga	g agctgccaac	ctttggccaa	gcccgctcat	gatcaaacgc	900
tctaagaaga acagcctgg	c cttgtccctg	acggccgacc	agatggtcag	tgccttgttg	960
gatgctgagc cccccatac	t ctattccgag	tatgatccta	ccagaccctt	cagtgaagct	1020
tcgatgatgg gcttactga	c caacctggca	gacagggagc	tggttcacat	gatcaactgg	1080

gcgaagaggg tgccaggctt tgtggatttg accetccatg atcaggtcca cettetagaa 1140

25136771 - 7 -

tgtgcctggc	tagagatcct	gatgattggt	ctcgtctggc	gctccatgga	gcacccagtg	1200
aagctactgt	ttgctcctaa	cttgctcttg	gacaggaacc	agggaaaatg	tgtagagggc	1260
atggtggaga	tcttcgacat	gctgctggct	acatcatctc	ggttccgcat	gatgaatctg	1320
cagggagagg	agtttgtgtg	cctcaaatct	attattttgc	ttaattctgg	agtgtacaca	1380
tttctgtcca	gcaccctgaa	gtctctggaa	gagaaggacc	atatccaccg	agtcctggac	1440
aagatcacag	acactttgat	ccacctgatg	gccaaggcag	gcctgaccct	gcagcagcag	1500
caccagcggc	tggcccagct	cctcctcatc	ctctcccaca	tcaggcacat	gaggaaccag	1560
ggaaaatgtg	tagagggcat	ggtggagatc	ttcgacatgc	tgctggctac	atcatctcgg	1620
ttccgcatga	tgaatctgca	gggagaggag	tttgtgtgcc	tcaaatctat	tattttgctt	1680
aattctggag	tgtacacatt	tctgtccagc	accctgaagt	ctctggaaga	gaaggaccat	1740
atccaccgag	tcctggacaa	gatcacagac	actttgatcc	acctgatggc	caaggcaggc	1800
ctgaccctgc	agcagcagca	ccagcggctg	gcccagctcc	tcctcatcct	ctcccacatc	1860
aggcacatga	gtaacaaagg	catggagcat	ctgtacagca	tgaagtgcaa	gaacgtggtg	1920
cccctctatg	acctgctgct	ggagatgctg	gacgcccacc	gcctacatgc	gcccactagc	1980
cgtggaggg	catccgtgga	ggagacggac	caaagccact	tggccactgc	gggctctact	2040
tcatcgcatt	ccttgcaaaa	gtattacatc	acgggggagg	cagagggttt	ccctgccaca	2100
tcatcgcatt gtctga	ccttgcaaaa	gtattacatc	acgggggagg	cagagggttt	ccctgccaca	2100 2106
	2	gtattacatc	acgggggagg	cagagggttt	ccctgccaca	
gtctga  <210> 6 <211> 2096 <212> DNA <213> Huma <400> 6	2					
gtctga  <210> 6 <211> 2096 <212> DNA <213> Huma <400> 6 gaattccaaa	2 an	ttcttgtatt	tttgatgaag	gagaaatact	gtaatgatca	2106
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac	2 an attgtgatgt	ttcttgtatt ttaggccagc	tttgatgaag cctttgtagc	gagaaatact gttatacaaa	gtaatgatca ctgaaagcac	2106
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg	2 an attgtgatgt tatgtacact	ttcttgtatt ttaggccagc gggcagggcc	tttgatgaag cctttgtagc ggggccagag	gagaaatact gttatacaaa ctcgcgtgtc	gtaatgatca ctgaaagcac ggcgggacat	2106 60 120
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg gcgctgcgtc	2 an attgtgatgt tatgtacact caggctcccg	ttcttgtatt ttaggccagc gggcagggcc tcgggctgtg	tttgatgaag cctttgtagc ggggccagag ctcttttcc	gagaaatact gttatacaaa ctcgcgtgtc aggtggcccg	gtaatgatca ctgaaagcac ggcgggacat ccggtttctg	2106 60 120 180
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg gcgctgcgtc agccttctgc	an attgtgatgt tatgtacact caggctcccg gcctctaacc	ttcttgtatt ttaggccagc gggcagggcc tcgggctgtg cacggtctgc	tttgatgaag cctttgtagc ggggccagag ctcttttcc accctgcccg	gagaaatact gttatacaaa ctcgcgtgtc aggtggcccg cggccacgga	gtaatgatca ctgaaagcac ggcgggacat ccggtttctg ccatgaccat	2106 60 120 180 240
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg gcgctgcgtc agccttctgc gaccctccac	an attgtgatgt tatgtacact caggctcccg gcctctaacc cctgcgggga	ttcttgtatt ttaggccagc gggcagggcc tcgggctgtg cacggtctgc ctgggatggc	tttgatgaag cctttgtagc ggggccagag ctcttttcc accctgcccg cctactgcat	gagaaatact gttatacaaa ctcgcgtgtc aggtggcccg cggccacgga cagatccaag	gtaatgatca ctgaaagcac ggcgggacat ccggtttctg ccatgaccat ggaacgagct	2106 60 120 180 240 300
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg gcgctgcgtc agccttctgc gaccctccac ggagccctg	an attgtgatgt tatgtacact caggctcccg gcctctaacc cctgcgggga accaaagcat	ttcttgtatt ttaggccagc gggcagggcc tcgggctgtg cacggtctgc ctgggatggc agctcaagat	tttgatgaag cctttgtagc ggggccagag ctcttttcc accctgcccg cctactgcat ccccctggag	gagaaatact gttatacaaa ctcgcgtgtc aggtggcccg cggccacgga cagatccaag	gtaatgatca ctgaaagcac ggcgggacat ccggtttctg ccatgaccat ggaacgagct gcgaggtgta	2106 60 120 180 240 300 360
gtctga  <210> 6 <211> 2093 <212> DNA <213> Huma <400> 6 gaattccaaa ctgtttacac accggacccg gcgctgcgtc agccttctgc gaccctccac ggagcccctg cctggacagc	an attgtgatgt tatgtacact caggctcccg gcctctaacc cctgcgggga accaaagcat aaccgtccgc	ttcttgtatt ttaggccagc gggcagggcc tcgggctgtg cacggtctgc ctgggatggc agctcaagat ccgtgtacaa	tttgatgaag cctttgtagc ggggccagag ctcttttcc accctgcccg cctactgcat ccccctggag ctaccccgag	gagaaatact gttatacaaa ctcgcgtgtc aggtggcccg cggccacgga cagatccaag cggcccctgg	gtaatgatca ctgaaagcac ggcgggacat ccggtttctg ccatgaccat ggaacgagct gcgaggtgta acgagttcaa	2106 60 120 180 240 300 360 420

gtctgaggct gcggcgttcg gctccaacgg cctggggggt ttccccccac tcaacagcgt

600

25136771 - 8 -

gtctccgagc	ccgctgatgc	tactgcaccc	gccgccgcag	ctgtcgcctt	tcctgcagcc	660
ccacggccag	caggtgccct	actacctgga	gaacgagccc	agcggctaca	cggtgcgcga	720
ggccggcccg	ccggcattct	acaggccaaa	ttcagataat	cgacgccagg	gtggcagaga	780
aagattggcc	agtaccaatg	acaagggaag	tatggctatg	gaatctgcca	aggagactcg	840
ctactgtgca	gtgtgcaatg	actatgcttc	aggctaccat	tatggagtct	ggtcctgtga	900
gggctgcaag	gccttcttca	agagaagtat	tcaaggacat	aacgactata	tgtgtccagc	960
caccaaccag	tgcaccattg	ataaaaacag	gaggaagagc	tgccaggcct	gccggctccg	1020
caaatgctac	gaagtgggaa	tgatgaaagg	tgggatacga	aaagaccgaa	gaggaggag	1080
aatgttgaaa	cacaagcgcc	agagagatga	tggggagggc	aggggtgaag	tggggtctgc	1140
tggagacatg	agagctgcca	acctttggcc	aagcccgctc	atgatcaaac	gctctaagaa	1200
gaacagcctg	gccttgtccc	tgacggccga	ccagatggtc	agtgccttgt	tggatgctga	1260
gcccccata	ctctattccg	agtatgatcc	taccagaccc	ttcagtgaag	cttcgatgat	1320
gggcttactg	accaacctgg	cagacaggga	gctggttcac	atgatcaact	gggcgaagag	1380
ggtgccaggc	tttgtggatt	tgaccctcca	tgatcaggtc	caccttctag	aatgtgcctg	1440
gctagagatc	ctgatgattg	gtctcgtctg	gcgctccatg	gagcacccag	tgaagctact	1500
gtttgctcct	aacttgctct	tggacaggaa	ccagggaaaa	tgtgtagagg	gcatggtgga	1560
gatcttcgac	atgctgctgg	ctacatcatc	tcggttccgc	atgatgaatc	tgcagggaga	1620
ggagtttgtg	tgcctcaaat	ctattatttt	gcttaattct	ggagtgtaca	catttctgtc	1680
cagcaccctg	aagtctctgg	aagagaagga	ccatatccac	cgagtcctgg	acaagatcac	1740
agacactttg	atccacctga	tggccaaggc	aggcctgacc	ctgcagcagc	agcaccagcg	1800
gctggcccag	ctcctcctca	tcctctccca	catcaggcac	atgagtaaca	aaggcatgga	1860
gcatctgtac	agcatgaagt	gcaagaacgt	ggtgcccctc	tatgacctgc	tgctggagat	1920
gctggacgcc	caccgcctac	atgcgcccac	tagccgtgga	ggggcatccg	tggaggagac	1980
ggaccaaagc	cacttggcca	ctgcgggctc	tacttcatcg	cattccttgc	aaaagtatta	2040
catcacgggg	gaggcagagg	gtttccctgc	cacagtctga	gagctccctg	gc	2092

<210> 7 <211> 6450 <212> DNA

<213> Human

<400> 7

gagttgtgcc tggagtgatg tttaagccaa tgtcagggca aggcaacagt ccctggccgt 60 cctccagcac ctttgtaatg catatgagct cgggagacca gtacttaaag ttggaggccc 120

25136771 - 9 -

gggagcccag	gagctggcgg	agggcgttcg	tcctgggagc	tgcacttgct	ccgtcgggtc	180
gccggcttca	ccggaccgca	ggctcccggg	gcagggccgg	ggccagagct	cgcgtgtcgg	240
cgggacatgc	gctgcgtcgc	ctctaacctc	gggctgtgct	ctttttccag	gtggcccgcc	300
ggtttctgag	ccttctgccc	tgcggggaca	cggtctgcac	cctgcccgcg	gccacggacc	360
atgaccatga	ccctccacac	caaagcatct	gggatggccc	tactgcatca	gatccaaggg	420
aacgagctgg	agcccctgaa	ccgtccgcag	ctcaagatcc	ccctggagcg	gcccctgggc	480
gaggtgtacc	tggacagcag	caagcccgcc	gtgtacaact	accccgaggg	cgccgcctac	540
gagttcaacg	ccgcggccgc	cgccaacgcg	caggtctacg	gtcagaccgg	cctcccctac	600
ggccccgggt	ctgaggctgc	ggcgttcggc	tccaacggcc	tggggggttt	cccccactc	660
aacagcgtgt	ctccgagccc	gctgatgcta	ctgcacccgc	cgccgcagct	gtcgcctttc	720
ctgcagcccc	acggccagca	ggtgccctac	tacctggaga	acgagcccag	cggctacacg	780
gtgcgcgagg	ccggcccgcc	ggcattctac	aggccaaatt	cagataatcg	acgccagggt	840
ggcagagaaa	gattggccag	taccaatgac	aagggaagta	tggctatgga	atctgccaag	900
gagactcgct	actgtgcagt	gtgcaatgac	tatgcttcag	gctaccatta	tggagtctgg	960
tcctgtgagg	gctgcaaggc	cttcttcaag	agaagtattc	aaggacataa	cgactatatg	1020
tgtccagcca	ccaaccagtg	caccattgat	aaaaacagga	ggaagagctg	ccaggcctgc	1080
cggctccgca	aatgctacga	agtgggaatg	atgaaaggtg	ggatacgaaa	agaccgaaga	1140
ggagggagaa	tgttgaaaca	caagcgccag	agagatgatg	gggagggcag	gggtgaagtg	1200
gggtctgctg	gagacatgag	agctgccaac	ctttggccaa	gcccgctcat	gatcaaacgc	1260
tctaagaaga	acagcctggc	cttgtccctg	acggccgacc	agatggtcag	tgccttgttg	1320
gatgctgagc	ccccatact	ctattccgag	tatgatccta	ccagaccctt	cagtgaagct	1380
tcgatgatgg	gcttactgac	caacctggca	gacagggagc	tggttcacat	gatcaactgg	1440
gcgaagaggg	tgccaggctt	tgtggatttg	accctccatg	atcaggtcca	ccttctagaa	1500
tgtgcctggc	tagagatcct	gatgattggt	ctcgtctggc	gctccatgga	gcacccagtg	1560
aagctactgt	ttgctcctaa	cttgctcttg	gacaggaacc	agggaaaatg	tgtagagggc	1620
atggtggaga	tcttcgacat	gctgctggct	acatcatctc	ggttccgcat	gatgaatctg	1680
cagggagagg	agtttgtgtg	cctcaaatct	attattttgc	ttaattctgg	agtgtacaca	1740
tttctgtcca	gcaccctgaa	gtctctggaa	gagaaggacc	atatccaccg	agtcctggac	1800
aagatcacag	acactttgat	ccacctgatg	gccaaggcag	gcctgaccct	gcagcagcag	1860
caccagcggc	tggcccagct	cctcctcatc	ctctcccaca	tcaggcacat	gagtaacaaa	1920

25136771 - 10 -

ggcatggagc atctgtacag catgaagtgc aagaacgtgg tgcccctcta tgacctgctg 1980 ctggagatgc tggacgccca ccgcctacat gcgcccacta gccgtggagg ggcatccgtg 2040 gaggagacgg accaaagcca cttggccact gcgggctcta cttcatcgca ttccttgcaa 2100 aagtattaca tcacggggga ggcagagggt ttccctgcca cagtctgaga gctccctggc 2160 teccaeaegg tteagataat eeetgetgea ttttaeeete ateatgeace aetttageea 2220 aattetgtet eetgeataca eteeggeatg cateeaacae caatggettt etagatgagt 2280 ggccattcat ttgcttgctc agttcttagt ggcacatctt ctgtcttctg ttgggaacag 2340 ccaaagggat tccaaggcta aatctttgta acagctctct ttcccccttg ctatgttact 2400 aagcgtgagg attcccgtag ctcttcacag ctgaactcag tctatgggtt ggggctcaga 2460 taactctgtg catttaagct acttgtagag acccaggcct ggagagtaga cattttgcct 2520 ctgataagca ctttttaaat ggctctaaga ataagccaca gcaaagaatt taaagtggct 2580 cctttaattg gtgacttgga gaaagctagg tcaagggttt attatagcac cctcttgtat 2640 tcctatggca atgcatcctt ttatgaaagt ggtacacctt aaagctttta tatgactgta 2700 gcagagtatc tggtgattgt caattcactt ccccctatag gaatacaagg ggccacacag 2760 ggaaggcaga tcccctagtt ggccaagact tattttaact tgatacactg cagattcaga 2820 gtgtcctgaa gctctgcctc tggctttccg gtcatgggtt ccagttaatt catgcctccc 2880 atggacctat ggagagcaac aagttgatct tagttaagtc tccctatatg agggataagt 2940 tcctgatttt tgtttttatt tttgtgttac aaaagaaagc cctccctccc tgaacttgca 3000 gtaaggtcag cttcaggacc tgttccagtg ggcactgtac ttggatcttc ccggcgtgtg 3060 tgtgccttac acaggggtga actgttcact gtggtgatgc atgatgaggg taaatggtag 3120 ttgaaaggag caggggccct ggtgttgcat ttagccctgg ggcatggagc tgaacagtac 3180 ttgtgcagga ttgttgtggc tactagagaa caagagggaa agtagggcag aaactggata 3240 cagttetgag cacagecaga ettgetcagg tggecetgea caggetgeag etacetagga 3300 acatteettg cagaccccge attgeetttg ggggtgeeet gggateeetg gggtagteea 3360 getettatte attteeeage gtggeeetgg ttggaagaag eagetgteaa gttgtagaea 3420 gctgtgttcc tacaattggc ccagcaccct ggggcacggg agaagggtgg ggaccgttgc 3480 tgtcactact caggctgact ggggcctggt cagattacgt atgcccttgg tggtttagag 3540 ataatccaaa atcagggttt ggtttgggga agaaaatcct cccccttcct cccccgcccc 3600 gttccctacc gcctccactc ctgccagctc atttccttca atttcctttg acctataggc 3660 taaaaaagaa aggctcattc cagccacagg gcagccttcc ctgggccttt gcttctctag 3720 cacaattatg ggttacttcc tttttcttaa caaaaaagaa tgtttgattt cctctgggtg 3780

25136771 - 11 -

accttattgt ctgtaattga aaccctattg agaggtgatg tctgtgttag ccaatgaccc 3840 aggtagctgc tcgggcttct cttggtatgt cttgtttgga aaagtggatt tcattcattt 3900 ctgattgtcc agttaagtga tcaccaaagg actgagaatc tgggagggca aaaaaaaaa 3960 aaaaagtttt tatgtgcact taaatttggg gacaatttta tgtatctgtg ttaaggatat 4020 gcttaagaac ataattettt tgttgetgtt tgtttaagaa gcaeettagt ttgtttaaga 4080 agcaccttat atagtataat atatattttt ttgaaattac attgcttgtt tatcagacaa 4140 ttgaatgtag taattetgtt etggatttaa tttgaetggg ttaacatgca aaaaccaagg 4200 aaaaatattt agtttttttt tttttttttg tatacttttc aagctacctt gtcatgtata 4260 cagtcattta tgcctaaagc ctggtgatta ttcatttaaa tgaagatcac atttcatatc 4320 aacttttgta tccacagtag acaaaatagc actaatccag atgcctattg ttggatattg 4380 aatgacagac aatcttatgt agcaaagatt atgcctgaaa aggaaaatta ttcagggcag 4440 ctaattttgc ttttaccaaa atatcagtag taatattttt ggacagtagc taatgggtca 4500 gtgggttctt tttaatgttt atacttagat tttcttttaa aaaaattaaa ataaaacaaa 4560 aaaaatttct aggactagac gatgtaatac cagctaaagc caaacaatta tacagtggaa 4620 ggttttacat tattcatcca atgtgtttct attcatgtta agatactact acatttgaag 4680 tgggcagaga acatcagatg attgaaatgt tcgcccaggg gtctccagca actttggaaa 4740 tctctttgta tttttacttg aagtgccact aatggacagc agatattttc tggctgatgt 4800 tggtattggg tgtaggaaca tgatttaaaa aaaaaactct tgcctctgct ttcccccact 4860 ctgaggcaag ttaaaatgta aaagatgtga tttatctggg gggctcaggt atggtgggga 4920 agtggattca ggaatctggg gaatggcaaa tatattaaga agagtattga aagtatttgg 4980 aggaaaatgg ttaattctgg gtgtgcacca aggttcagta gagtccactt ctgccctgga 5040 gaccacaaat caactagctc catttacagc catttctaaa atggcagctt cagttctaga 5100 gaagaaagaa caacatcagc agtaaagtcc atggaatagc tagtggtctg tgtttctttt 5160 cgccattgcc tagcttgccg taatgattct ataatgccat catgcagcaa ttatgagagg 5220 ctaggtcatc caaagagaag accctatcaa tgtaggttgc aaaatctaac ccctaaggaa 5280 gtgcagtctt tgatttgatt tccctagtaa ccttgcagat atgtttaacc aagccatagc 5340 ccatgccttt tgagggctga acaaataagg gacttactga taatttactt ttgatcacat 5400 taaggtgttc tcaccttgaa atcttataca ctgaaatggc cattgattta ggccactggc 5460 ttagagtact cetteceetg catgacactg attacaaata ettteetatt cataetttee 5520 aattatgaga tggactgtgg gtactgggag tgatcactaa caccatagta atgtctaata 5580

25136771 - 12 -

ttcacag	gca gatctgcttg	gggaagctag	ttatgtgaaa	ggcaaataaa	gtcatacagt	5640
agctcaa	aag gcaaccataa	ttctctttgg	tgcaagtctt	gggagcgtga	tctagattac	5700
actgcac	cat tcccaagtta	atcccctgaa	aacttactct	caactggagc	aaatgaactt	5760
tggtccc	aaa tatccatctt	ttcagtagcg	ttaattatgc	tctgtttcca	actgcatttc	5820
ctttcca	att gaattaaagt	gtggcctcgt	ttttagtcat	ttaaaattgt	tttctaagta	5880
attgctg	cct ctattatggc	acttcaattt	tgcactgtct	tttgagattc	aagaaaaatt	5940
tctattc	att tttttgcatc	caattgtgcc	tgaactttta	aaatatgtaa	atgctgccat	6000
gttccaa	acc catcgtcagt	gtgtgtgttt	agagctgtgc	accctagaaa	caacatactt	6060
gtcccat	gag caggtgcctg	agacacagac	ccctttgcat	tcacagagag	gtcattggtt	6120
atagaga	ctt gaattaataa	gtgacattat	gccagtttct	gttctctcac	aggtgataaa	6180
caatgct	ttt tgtgcactac	atactcttca	gtgtagagct	cttgttttat	gggaaaaggc	6240
tcaaatg	cca aattgtgttt	gatggattaa	tatgcccttt	tgccgatgca	tactattact	6300
gatgtga	ctc ggttttgtcg	cagctttgct	ttgtttaatg	aaacacactt	gtaaacctct	6360
tttgcac	ttt gaaaaagaat	ccagcgggat	gctcgagcac	ctgtaaacaa	ttttctcaac	6420
ctatttç	gatg ttcaaataaa	gaattaaact				6450
<210>	8					
<211> <212>	7 DNA					
	Artificial Sec	[uence				
<220>	E-turnen Dogne	ngo Flomont				
<223>	Estrogen Respo	Mise Element				
<220>						
<221> <222>	misc_feature (4)(4)					
	N equals C or	G				
<400>						7
tgantc	a					,
<210>	9					
<211>	595					
<212>	PRT					
<213>	Human					

Gln Ile Gln Gly Asn Glu Leu Glu Pro Leu Asn Arg Pro Gln Leu Lys

25136771 - 13 -

<400> 9

20 25 30

Ile Pro Leu Glu Arg Pro Leu Gly Glu Val Tyr Leu Asp Ser Ser Lys  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

- Pro Ala Val Tyr Asn Tyr Pro Glu Gly Ala Ala Tyr Glu Phe Asn Ala 50 55 60
- Ala Ala Ala Asn Ala Gln Val Tyr Gly Gln Thr Gly Leu Pro Tyr 65 70 75 80
- Gly Pro Gly Ser Glu Ala Ala Ala Phe Gly Ser Asn Gly Leu Gly Gly 85 90 95
- Phe Pro Pro Leu Asn Ser Val Ser Pro Ser Pro Leu Met Leu His 100 105 110
- Pro Pro Gln Leu Ser Pro Phe Leu Gln Pro His Gly Gln Gln Val 115 120 125
- Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Gly Tyr Thr Val Arg Glu Ala 130 135 140
- Gly Pro Pro Ala Phe Tyr Arg Pro Asn Ser Asp Asn Arg Arg Gln Gly
  145 150 155 160
- Gly Arg Glu Arg Leu Ala Ser Thr Asn Asp Lys Gly Ser Met Ala Met 165 170 175
- Glu Ser Ala Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala 180 185 190
- Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe 195 200 205
- Phe Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr 210 215 220
- Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 225 230 235 240
- Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg 245 250 255
- Lys Asp Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 260 265 270

25136771 - 14 -

Asp Gly Glu Gly Arg Gly Glu Val Gly Ser Ala Gly Asp Met Arg Ala 280 Ala Asn Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn 295 300 Ser Leu Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 310 315 320 Asp Ala Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 330 325 Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 340 Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Val 355 360 Asp Leu Thr Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 375 370 Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val 390 Lys Leu Leu Phe Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys 415 405 Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 420 425 Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 450 Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 465 Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr Leu Gln Gln His Gln Arg Leu Ala Gln Leu Leu Ile Leu Ser

505

510

25136771 - 15 -

His Ile Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Ser Met 515

Lys Cys Lys Asn Val Val Pro Leu Tyr Asp Leu Leu Glu Met Leu 530 535

Asp Ala His Arg Leu His Ala Pro Thr Ser Arg Gly Gly Ala Ser Val 545 550

Glu Glu Thr Asp Gln Ser His Leu Ala Thr Ala Gly Ser Thr Ser Ser 565

His Ser Leu Gln Lys Tyr Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro 585

Ala Thr Val 595

<210> 10

<211> 547 <212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Estrogen Receptor Ligand Binding Domain Fusion

Met Val Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu 5

Val Glu Leu Asp Gly Asp Val Asn Gly His Arg Phe Ser Val Ser Gly 20 25

Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile 35 40

Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr 50 55

Leu Thr Trp Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys 70 7.5

Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu

Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu 100 105

25136771 - 16 - Val Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly 115 120 125

Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr 130 135 140

Asn Tyr Ile Ser His Asn Val Tyr Ile Thr Ala Asp Lys Gln Lys Asn 145 150 155 160

Gly Ile Lys Ala His Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser 165 170 175

Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly 180 185 190

Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu 195 200 205

Ser Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe 210 220

Val Thr Ala Ala Arg Met Pro Gly Gly Ser Ala Gly Asp Met Arg Ala 225 230 235 240

Ala Asn Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn 245 250 255

Ser Leu Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 260 265 270

Asp Ala Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 275 280 285

Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 290 295 300

Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Val 305 310 315 320

Asp Leu Thr Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 325 330 335

Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val 340 345 350

25136771 - 17 -

Lys Leu Leu Phe Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys 355 360 365

Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 370 380

Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu 385 390 395 400

Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 405 410 415

Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 420 425 430

Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 435 440 445

Leu Gln Gln Gln His Gln Arg Leu Ala Gln Leu Leu Leu Ile Leu Ser 450 455 460

His Ile Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Ser Met 465 470 475 480

Lys Cys Lys Asn Val Val Pro Leu Tyr Asp Leu Leu Glu Met Leu 485 490 495

Asp Ala His Arg Leu His Ala Pro Thr Ser Arg Gly Gly Ala Ser Val 500 505 510

Glu Glu Thr Asp Gln Ser His Leu Ala Thr Ala Gly Ser Thr Ser Ser 515 520 525

His Ser Leu Gln Lys Tyr Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro  $530 \hspace{1.5cm} 535 \hspace{1.5cm} 540$ 

Ala Thr Val 545

<210> 11

<211> 113

<212> PRT

<213> human

<400> 11

Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp Asp Gly

25136771 - 18 -

1 5 10 15

Glu Gly Arg Gly Glu Val Gly Ser Ala Gly Asp Met Arg Ala Ala Asn 20 25 30

Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn Ser Leu 35 40 45

Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu Asp Ala 50 55 60

Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro Phe Ser 65 70 75 80

Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg Glu Leu 85 90 95

Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Thr Arg Glu Asn 100 105 110

Val

<210> 12

<211> 701

<212> PRT

<213> human

<400> 12

Met Thr Met Thr Leu His Thr Lys Ala Ser Gly Met Ala Leu Leu His 1 5 10 15

Gln Ile Gln Gly Asn Glu Leu Glu Pro Leu Asn Arg Pro Gln Leu Lys 20 25 30

Ile Pro Leu Glu Arg Pro Leu Gly Glu Val Tyr Leu Asp Ser Ser Lys 35 40 45

Pro Ala Val Tyr Asn Tyr Pro Glu Gly Ala Ala Tyr Glu Phe Asn Ala 50 60

Ala Ala Ala Asn Ala Gln Val Tyr Gly Gln Thr Gly Leu Pro Tyr 65 70 75 80

Gly Pro Gly Ser Glu Ala Ala Ala Phe Gly Ser Asn Gly Leu Gly Gly 85 90 95

25136771 - 19 -

Phe Pro Pro Leu Asn Ser Val Ser Pro Ser Pro Leu Met Leu Leu His
100 105 110

Pro Pro Gln Leu Ser Pro Phe Leu Gln Pro His Gly Gln Gln Val 115 120 125

Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Gly Tyr Thr Val Arg Glu Ala 130 140

Gly Pro Pro Ala Phe Tyr Arg Pro Asn Ser Asp Asn Arg Arg Gln Gly
145 150 155 160

Gly Arg Glu Arg Leu Ala Ser Thr Asn Asp Lys Gly Ser Met Ala Met 165 170 175

Glu Ser Ala Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala 180 185 190

Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe 195 200 205

Phe Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr 210 215 220

Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 225 230 235 240

Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg 245 250 255

Lys Asp Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 260 265 270

Asp Gly Glu Gly Arg Gly Glu Val Gly Ser Ala Gly Asp Met Arg Ala 275 280 285

Ala Asn Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn 290 295 300

Ser Leu Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 305 310 315 320

Asp Ala Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 325 330 335

25136771

Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 340 Leu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Val 355

Asp Leu Thr Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 370 380

Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val 385 390 395 400

Lys Leu Leu Phe Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys 405 410 415

Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 420 425 430

Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu 435 440 445

Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 450 455 460

Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 465 470 475 480

Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 485 490 495

Leu Gln Gln His Gln Arg Leu Ala Gln Leu Leu Ile Leu Ser 500 505 510

His Ile Arg His Met Arg Asn Gln Gly Lys Cys Val Glu Gly Met Val 515 520 525

Glu Ile Phe Asp Met Leu Leu Ala Thr Ser Ser Arg Phe Arg Met Met 530 540

Asn Leu Gln Gly Glu Glu Phe Val Cys Leu Lys Ser Ile Ile Leu Leu 545 550 555 560

Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser Thr Leu Lys Ser Leu Glu 565 570 575

Glu Lys Asp His Ile His Arg Val Leu Asp Lys Ile Thr Asp Thr Leu

25136771

Ile	His	Leu 595	Met	Ala	Lys	Ala	Gly 600	Leu	Thr	Leu	Gln	Gln 605	Gln	His	Gln
Arg	Leu 610	Ala	Gln	Leu	Leu	Leu 615	Ile	Leu	Ser	His	Ile 620	Arg	His	Met	Ser

585

590

Asn Lys Gly Met Glu His Leu Tyr Ser Met Lys Cys Lys Asn Val Val 625 630 635 640

Pro Leu Tyr Asp Leu Leu Leu Glu Met Leu Asp Ala His Arg Leu His 645 650 655

Ala Pro Thr Ser Arg Gly Gly Ala Ser Val Glu Glu Thr Asp Gln Ser 660 665 670

His Leu Ala Thr Ala Gly Ser Thr Ser Ser His Ser Leu Gln Lys Tyr  $675 \hspace{1.5cm} 680 \hspace{1.5cm} 685$ 

Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro Ala Thr Val 690 695 700

<210> 13

<211> 595

<212> PRT

<213> human

580

<400> 13

Met Thr Met Thr Leu His Thr Lys Ala Ser Gly Met Ala Leu Leu His 1 5 10 15

Gln Ile Gln Gly Asn Glu Leu Glu Pro Leu Asn Arg Pro Gln Leu Lys 20 25 30

Ile Pro Leu Glu Arg Pro Leu Gly Glu Val Tyr Leu Asp Ser Ser Lys 35 40 45

Pro Ala Val Tyr Asn Tyr Pro Glu Gly Ala Ala Tyr Glu Phe Asn Ala 50 55 60

Ala Ala Ala Asn Ala Gln Val Tyr Gly Gln Thr Gly Leu Pro Tyr 65 70 75 80

Gly Pro Gly Ser Glu Ala Ala Phe Gly Ser Asn Gly Leu Gly Gly 85 90 95

25136771 - 22 -

Phe Pro Pro Leu Asn Ser Val Ser Pro Ser Pro Leu Met Leu His
100 105 110

Pro Pro Pro Gln Leu Ser Pro Phe Leu Gln Pro His Gly Gln Gln Val 115 120 125

Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Gly Tyr Thr Val Arg Glu Ala 130 135 140

Gly Pro Pro Ala Phe Tyr Arg Pro Asn Ser Asp Asn Arg Arg Gln Gly 145 150 155 160

Gly Arg Glu Arg Leu Ala Ser Thr Asn Asp Lys Gly Ser Met Ala Met 165 170 175

Glu Ser Ala Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala 180 185 190

Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe 195 200 205

Phe Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr 210 215 220

Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 225 230 235 240

Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg 245 250 255

Lys Asp Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 260 265 270

Asp Gly Glu Gly Arg Gly Glu Val Gly Ser Ala Gly Asp Met Arg Ala 275 280 285

Ala Asn Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn 290 295 300

Ser Leu Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 305 310 315 320

Asp Ala Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 325 330 335

25136771 - 23 -

Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 345 340 Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Val 355 360 Asp Leu Thr Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 370 375 380 Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val 395 Lys Leu Leu Phe Ala Pro Asn Leu Leu Asp Arg Asn Gln Gly Lys Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 420 Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Phe Val Cys Leu 435 Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 450 Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 470 Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 485 490 Leu Gln Gln His Gln Arg Leu Ala Gln Leu Leu Ile Leu Ser 500 His Ile Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Ser Met 515 520 Lys Cys Lys Asn Val Val Pro Leu Tyr Asp Leu Leu Glu Met Leu 530 Asp Ala His Arg Leu His Ala Pro Thr Ser Arg Gly Gly Ala Ser Val 545 550 560 Glu Glu Thr Asp Gln Ser His Leu Ala Thr Ala Gly Ser Thr Ser Ser 565

His Ser Leu Gln Lys Tyr Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro

25136771 - 24 -

580 585 590

Ala Thr Val 595

<210> 14

<211> 595

<212> PRT

<213> Human

<400> 14

Gln Ile Gln Gly Asn Glu Leu Glu Pro Leu Asn Arg Pro Gln Leu Lys 20 25 30

Ile Pro Leu Glu Arg Pro Leu Gly Glu Val Tyr Leu Asp Ser Ser Lys 35 40 45

Pro Ala Val Tyr Asn Tyr Pro Glu Gly Ala Ala Tyr Glu Phe Asn Ala 50 55 60

Ala Ala Ala Asn Ala Gln Val Tyr Gly Gln Thr Gly Leu Pro Tyr 65 70 75 80

Gly Pro Gly Ser Glu Ala Ala Ala Phe Gly Ser Asn Gly Leu Gly Gly 85 90 95

Pro Pro Gln Leu Ser Pro Phe Leu Gln Pro His Gly Gln Gln Val

Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Gly Tyr Thr Val Arg Glu Ala 130 135 140

Gly Pro Pro Ala Phe Tyr Arg Pro Asn Ser Asp Asn Arg Arg Gln Gly 145 150 155 160

Gly Arg Glu Arg Leu Ala Ser Thr Asn Asp Lys Gly Ser Met Ala Met 165 170 175

Glu Ser Ala Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala . 180 185 190 Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe 195 200 205

Phe Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr 210 215 220

Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 225 230 235 240

Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg 245 250 255

Lys Asp Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 260 265 270

Asp Gly Glu Gly Arg Gly Glu Val Gly Ser Ala Gly Asp Met Arg Ala 275 280 285

Ala Asn Leu Trp Pro Ser Pro Leu Met Ile Lys Arg Ser Lys Lys Asn 290 295 300

Ser Leu Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 305 310 315 320

Asp Ala Glu Pro Pro Ile Leu Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 325 330 335

Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 340 345 350

Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Val 355 360 365

Asp Leu Thr Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 370 380

Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Val 385 390 395 400

Lys Leu Leu Phe Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys 405 410 415

Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 420 425 430

25136771 - 26 -

Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu 435 440 445

Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 450 455 460

Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 465 470 480

Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 485 490 495

Leu Gln Gln His Gln Arg Leu Ala Gln Leu Leu Leu Ile Leu Ser 500 505 510

His Ile Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Ser Met 515 520 525

Lys Cys Lys Asn Val Val Pro Leu Tyr Asp Leu Leu Leu Glu Met Leu 530 540

Asp Ala His Arg Leu His Ala Pro Thr Ser Arg Gly Gly Ala Ser Val 545 550 555 560

Glu Glu Thr Asp Gln Ser His Leu Ala Thr Ala Gly Ser Thr Ser Ser 565 570 575

His Ser Leu Gln Lys Tyr Tyr Ile Thr Gly Glu Ala Glu Gly Phe Pro 580 585 590

Ala Thr Val 595

<210> 15

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 15

caagcgccag agagatgatg

20

<210> 16

<211> 20

<212> DNA

<213> Artificial Sequence

25136771

<220> <223> Primer		
<400> 16		
acaaggcact gaccatctgg		20
<210> 17 <211> 20 <212> DNA <213> Artificial Sequence		
<220> <223> Primer		
<400> 17		
gaccatctgg tcggccgtca		20
		20
<210> 18 <211> 21		
<212> DNA		
<213> Artificial Sequence		
<220> <223> Primer		
<400> 18		
cagagagaat gatggggagg g		21
		2.1
<210> 19 <211> 11522		
<211> 11322 <212> DNA		
<213> Artificial Sequence		
<220>		
<223> Vector pER8		
<pre>&lt;400&gt; 19 gtttacccgc caatatatcc tgtcaaacac tgatagttt</pre>	2.2200000000000000000000000000000000000	
		60
aatctgatcc aagctcaagc taagcttgca tgcctgcag		120
ccacgtgccg ccacgtgccg ccacgtgcc		180
gacgtaaggg atgacgcaca atcccactat ccttcgcaa	g accettecte tatataagga	240
agttcatttc atttggagag gacacgctgg gatccccaa	t tccgggcgga atgaaagcgt	300
taacggccag gcaacaagag gtgtttgatc tcatccgtga	a tcacatcagc cagacaggta	360
tgccgccgac gcgtgcggaa atcgcgcagc gtttggggtt		420
aagaacatct gaaggcgctg gcacgcaaag gcgttattga	a aattgtttcc ggcgcatcac	480
gcgggattcg tctgttgcag gaagaggaag aagggttgcc		540
ccggtgaacc gtcgagcgcc cccccgaccg atgtcagcct		600
acggcgagga cgtggcgatg gcgcatgccg acgcgctaga		660

25136771

tgggggacgg	ggattccccg	ggtccgggat	ttacccccca	cgactccgcc	ccctacggcg	720
ctctggatat	ggccgacttc	gagtttgagc	agatgtttac	cgatgccctt	ggaattgacg	780
agtacggtgg	ggatccgtct	gctggagaca	tgagagctgc	caacctttgg	ccaagcccgc	840
tcatgatcaa	acgctctaag	aagaacagcc	tggccttgtc	cctgacggcc	gaccagatgg	900
tcagtgcctt	gttggatgct	gagcccccca	tactctattc	cgagtatgat	cctaccagac	960
ccttcagtga	agcttcgatg	atgggcttac	tgaccaacct	ggcagacagg	gagctggttc	1020
acatgatcaa	ctgggcgaag	agggtgccag	gctttgtgga	tttgaccctc	catgatcagg	1080
tccaccttct	agaatgtgcc	tggctagaga	tcctgatgat	tggtctcgtc	tggcgctcca	1140
tggagcaccc	agtgaagcta	ctgtttgctc	ctaacttgct	cttggacagg	aaccagggaa	1200
aatgtgtaga	gggcatggtg	gagatcttcg	acatgctgct	ggctacatca	tctcggttcc	1260
gcatgatgaa	tctgcaggga	gaggagtttg	tgtgcctcaa	atctattatt	ttgcttaatt	1320
ctggagtgta	cacatttctg	tccagcaccc	tgaagtctct	ggaagagaag	gaccatatcc	1380
accgagtcct	ggacaagatc	acagacactt	tgatccacct	gatggccaag	gcaggcctga	1440
ccctgcagca	gcagcaccag	cggctggccc	agctcctcct	catcctctcc	cacatcaggc	1500
acatgagtaa	caaaggcatg	gagcatctgt	acagcatgaa	gtgcaagaac	gtggtgcccc	1560
tctatgacct	gctgctggag	atgctggacg	cccaccgcct	acatgcgccc	actagccgtg	1620
gaggggcatc	cgtggaggag	acggaccaaa	gccacttggc	cactgcgggc	tctacttcat	1680
cgcattcctt	gcaaaagtat	tacatcacgg	gggaggcaga	gggtttccct	gccacagtct	1740
gagageteee	tggcgaattc	ccagagatgt	tagctgaaat	catcactaat	cagataccaa	1800
aatattcaaa	tggaaatatc	aaaaagcttc	tgtttcatca	aaaatgactc	gacctaactg	1860
agtaagctag	cttgttcgag	tattatggca	ttgggaaaac	tgtttttctt	gtaccatttg	1920
ttgtgcttgt	aatttactgt	gttttttatt	cggttttcgc	tatcgaactg	tgaaatggaa	1980
atggatggag	aagagttaat	gaatgatatg	gtccttttgt	tcattctcaa	attaatatta	2040
tttgttttt	ctcttatttg	ttgtgtgttg	aatttgaaat	tataagagat	atgcaaacat	2100
tttgttttga	gtaaaaatgt	gtcaaatcgt	ggcctctaat	gaccgaagtt	aatatgagga	2160
gtaaaacatc	ccaaacaagc	ttggaaactg	aaggcgggaa	acgacaatct	gatcatgagc	2220
ggagaattaa	gggagtcacg	ttatgacccc	cgccgatgac	gcgggacaag	ccgttttacg	2280
tttggaactg	acagaaccgc	aacgattgaa	ggagccactc	agccgcgggt	ttctggagtt	2340
taatgagcta	agcacatacg	tcagaaacca	ttattgcgcg	ttcaaaagtc	gcctaaggtc	2400
actatcagct	agcaaatatt	tcttgtcaaa	aatgctccac	tgacgttcca	taaattcccc	2460

25136771 - 29 -

tcggtatcca attagagtct catattcact ctcaatccaa ataatctgca ccggatcccc 2520 tagaatgaaa aagcctgaac tcaccgcgac gtctgtcgag aagtttctga tcgaaaagtt 2580 cgacagcgtc tccgacctga tgcagctctc ggagggcgaa gaatctcgtg ctttcagctt 2640 cgatgtagga gggcgtggat atgtcctgcg ggtaaatagc tgcgccgatg gtttctacaa 2700 agatcgttat gtttatcggc actttgcatc ggccgcgctc ccgattccgg aagtgcttga 2760 cattggggaa ttcagcgaga gcctgaccta ttgcatctcc cgccgtgcac agggtgtcac 2820 gttgcaagac ctgcctgaaa ccgaactgcc cgctgttctg cagccggtcg cggaggccat 2880 ggatgcgatc gctgcggccg atcttagcca gacgagcggg ttcggcccat tcggaccgca 2940 aggaatcggt caatacacta catggcgtga tttcatatgc gcgattgctg atccccatgt 3000 gtatcactgg caaactgtga tggacgacac cgtcagtgcg tccgtcgcgc aggctctcga 3060 tgagctgatg ctttgggccg aggactgccc cgaagtccgg cacctcgtgc acgcggattt 3120 cggctccaac aatgtcctga cggacaatgg ccgcataaca gcggtcattg actggagcga 3180 ggcgatgttc ggggattccc aatacgaggt cgccaacatc ttcttctgga ggccgtggtt 3240 ggcttgtatg gagcagcaga cgcgctactt cgagcggagg catccggagc ttgcaggatc 3300 gccgcggctc cgggcgtata tgctccgcat tggtcttgac caactctatc agagcttggt 3360 tgacggcaat ttcgatgatg cagettggge gcagggtega tgegaegeaa tegteegate 3420 cggagccggg actgtcgggc gtacacaaat cgcccgcaga agcgcggccg tctggaccga 3480 tggctgtgta gaagtactcg ccgatagtgg aaaccgacgc cccagcactc gtccgagggc 3540 aaaggaatag cgatcgttca aacatttggc aataaagttt cttaagattg aatcctgttg 3600 ccggtcttgc gatgattatc atataatttc tgttgaatta cgttaagcat gtaataatta 3660 acatgtaatg catgacgtta tttatgagat gggtttttat gattagagtc ccgcaattat 3720 acatttaata cgcgatagaa aacaaaatat agcgcgcaaa ctaggataaa ttatcgcgcg 3780 cggtgtcatc tatgttacta gatcggggaa ttgatccccc ctcgacagct tgcatgccag 3840 cttgggctgc aggtcgaggc taaaaaacta atcgcattat catcccctcg acgtactgta 3900 catataacca ctggttttat atacagcagt actgtacata taaccactgg ttttatatac 3960 agcagtcgac gtactgtaca tataaccact ggttttatat acagcagtac tgtacatata 4020 accactggtt ttatatacag cagtcgaggt aagattagat atggatatgt atatggatat 4080 gtatatggtg gtaatgccat gtaatatgct cgactctagg atcttcgcaa gacccttcct 4140 ctatataagg aagttcattt catttggaga ggacacgctg aagctagtcg actctagcct 4200 cgaggcgcgc cgggcccagg cctacgcgtt taattaacta gtcgatccag gcctcccagc 4260 tttcgtccgt atcatcggtt tcgacaacgt tcgtcaagtt caatgcatca gtttcattgc 4320

25136771 - 30 -

4380 ccacacca gaatcctact aagtttgagt attatggcat tggaaaagct gttttcttct atcatttgtt ctgcttgtaa tttactgtgt tctttcagtt tttgttttcg gacatcaaaa 4440 4500 tgcaaatgga tggataagag ttaataaatg atatggtcct tttgttcatt ctcaaattat 4560 tattatctgt tgtttttact ttaatgggtt gaatttaagt aagaaaggaa ctaacagtgt gatattaagg tgcaatgtta gacatataaa acagtctttc acctctcttt ggttatgtct 4620 tgaattggtt tgtttcttca cttatctgtg taatcaagtt tactatgagt ctatgatcaa 4680 4740 gtaattatgc aatcaagtta agtacagtat aggctttttg tgtcgagggg gtaccgagtc 4800 gaggaattca ctggccgtcg ttttacaacg tcgtgactgg gaaaaccctg gcgttaccca acttaatcgc cttgcagcac atcccccttt cgccagctgg cgtaatagcg aagaggcccg 4860 4920 caccgatcgc ccttcccaac agttgcgcag cctgaatggc gggtaccgag ctcgaattca atteggegtt aatteagtae attaaaaaeg teegeaatgt gttattaagt tgtetaageg 4980 tcaatttgtt tacaccacaa tatatcctgc caccagccag ccaacagctc cccgaccggc 5040 5100 ageteggeae aaaateaeea etegataeag geageeeate agteegggae ggegteageg ggagagccgt tgtaaggcgg cagactttgc tcatgttacc gatgctattc ggaagaacgg 5160 5220 caactaaget geegggtttg aaacaeggat gatetegegg agggtageat gttgattgta 5280 acgatgacag agcgttgctg cctgtgatca attcgggcac gaacccagtg gacataagcc 5340 tegtteggtt egtaagetgt aatgeaagta gegtaaetge egteaegeaa etggteeaga 5400 accttgaccg aacgcagcgg tggtaacggc gcagtggcgg ttttcatggc ttcttgttat 5460 gacatgtttt tttggggtac agtctatgcc tcgggcatcc aagcagcaag cgcgttacgc 5520 cgtgggtcga tgtttgatgt tatggagcag caacgatgtt acgcagcagg gcagtcgcc 5580 taaaacaaag ttaaacatca tgggggaagc ggtgatcgcc gaagtatcga ctcaactatc 5640 agaggtagtt ggcgtcatcg agcgccatct cgaaccgacg ttgctggccg tacatttgta 5700 cggctccgca gtggatggcg gcctgaagcc acacagtgat attgatttgc tggttacggt 5760 gaccgtaagg cttgatgaaa caacgcggcg agctttgatc aacgaccttt tggaaacttc ggcttcccct ggagagagcg agattctccg cgctgtagaa gtcaccattg ttgtgcacga 5820 cgacatcatt ccgtggcgtt atccagctaa gcgcgaactg caatttggag aatggcagcg 5880 caatgacatt cttgcaggta tcttcgagcc agccacgatc gacattgatc tggctatctt 5940 6000 gctgacaaaa gcaagagaac atagcgttgc cttggtaggt ccagcggcgg aggaactctt tgatccggtt cctgaacagg atctatttga ggcgctaaat gaaaccttaa cgctatggaa 6060 ctcgccgccc gactgggctg gcgatgagcg aaatgtagtg cttacgttgt cccgcatttg 6120

25136771 - 31 -

gtacagegea gtaaceggea aaategegee gaaggatgte getgeegaet gggeaatgga 6180 gcgcctgccg gcccagtatc agcccgtcat acttgaagct agacaggctt atcttggaca 6240 agaagaagat cgcttggcct cgcgcgcaga tcagttggaa gaatttgtcc actacgtgaa 6300 aggcgagatc accaaggtag tcggcaaata atgtctagct agaaattcgt tcaagccgac 6360 gccgcttcgc cggcgttaac tcaagcgatt agatgcacta agcacataat tgctcacagc 6420 caaactatca ggtcaagtct gcttttatta tttttaagcg tgcataataa gccctacaca 6480 aattgggaga tatatcatgc atgaccaaaa tcccttaacg tgagttttcg ttccactgag 6540 cgtcagaccc cgtagaaaag atcaaaggat cttcttgaga tccttttttt ctgcgcgtaa 6600 tctgctgctt gcaaacaaaa aaaccaccgc taccagcggt ggtttgtttg ccggatcaag 6660 agctaccaac tettttteeg aaggtaactg getteageag agegeagata ccaaatactg 6720 teettetagt gtageegtag ttaggeeace aetteaagaa etetgtagea eegeetaeat 6780 acctcgctct gctaatcctg ttaccagtgg ctgctgccag tggcgataag tcgtgtctta 6840 ccgggttgga ctcaagacga tagttaccgg ataaggcgca gcggtcgggc tgaacggggg 6900 gttcgtgcac acageccage ttggagegaa egaectacae egaaetgaga tacetacage 6960 gtgagctatg agaaagcgcc acgcttcccg aagggagaaa ggcggacagg tatccggtaa 7020 gcggcagggt cggaacagga gagcgcacga gggagcttcc agggggaaac gcctggtatc 7080 tttatagtcc tgtcgggttt cgccacctct gacttgagcg tcgatttttg tgatgctcgt 7140 caggggggcg gagcctatgg aaaaacgcca gcaacgcggc ctttttacgg ttcctggcct 7200 tttgctggcc ttttgctcac atgttctttc ctgcgttatc ccctgattct gtggataacc 7260 gtattaccgc ctttgagtga gctgataccg ctcgccgcag ccgaacgacc gagcgcagcg 7320 agtcagtgag cgaggaagcg gaagagcgcc tgatgcggta ttttctcctt acgcatctgt 7380 gcggtatttc acaccgcata tggtgcactc tcagtacaat ctgctctgat gccgcatagt 7440 taagccagta tacactccgc tatcgctacg tgactgggtc atggctgcgc cccgacaccc 7500 gccaacaccc gctgacgcgc cctgacgggc ttgtctgctc ccggcatccg cttacagaca 7560 agctgtgacc gtctccggga gctgcatgtg tcagaggttt tcaccgtcat caccgaaacg 7620 cgcgaggcag ggtgccttga tgtgggcgcc ggcggtcgag tggcgacggc gcggcttgtc 7680 cgcgccctgg tagattgcct ggccgtaggc cagccatttt tgagcggcca gcggccgcga 7740 taggccgacg cgaagcggcg gggcgtaggg agcgcagcga ccgaagggta ggcgcttttt 7800 gcagctcttc ggctgtgcgc tggccagaca gttatgcaca ggccaggcgg gttttaagag 7860 ttttaataag ttttaaagag ttttaggcgg aaaaatcgcc ttttttctct tttatatcag 7920 tcacttacat gtgtgaccgg ttcccaatgt acggctttgg gttcccaatg tacgggttcc 7980

8040 ggttcccaat gtacggcttt gggttcccaa tgtacgtgct atccacagga aagagacctt 8100 ttcgaccttt ttcccctgct agggcaattt gccctagcat ctgctccgta cattaggaac 8160 cggcggatgc ttcgccctcg atcaggttgc ggtagcgcat gactaggatc gggccagcct gccccgcctc ctccttcaaa tcgtactccg gcaggtcatt tgacccgatc agcttgcgca 8220 cggtgaaaca gaacttcttg aactctccgg cgctgccact gcgttcgtag atcgtcttga 8280 acaaccatct ggcttctgcc ttgcctgcgg cgcggcgtgc caggcggtag agaaaacggc 8340 cgatgccggg atcgatcaaa aagtaatcgg ggtgaaccgt cagcacgtcc gggttcttgc 8400 8460 cttctgtgat ctcgcggtac atccaatcag ctagctcgat ctcgatgtac tccggccgcc 8520 cggtttcgct ctttacgatc ttgtagcggc taatcaaggc ttcaccctcg gataccgtca ccaggeggee gttettggee ttettegtae getgeatgge aacgtgegtg gtgtttaace 8580 8640 gaatgcaggt ttctaccagg tcgtctttct gctttccgcc atcggctcgc cggcagaact 8700 8760 ggtatcggtt catggattcg gttagatggg aaaccgccat cagtaccagg tcgtaatccc 8820 acacactggc catgeeggee ggeeetgegg aaacetetae gtgeeegtet ggaagetegt ageggateae etegeeaget egteggteae gettegaeag aeggaaaaeg geeaegteea 8880 8940 tgatgctgcg actatcgcgg gtgcccacgt catagagcat cggaacgaaa aaatctggtt 9000 gctcgtcgcc cttgggcggc ttcctaatcg acggcgcacc ggctgccggc ggttgccggg 9060 attetttgeg gattegatea geggeegett geeaegatte accggggegt gettetgeet cgatgcgttg ccgctgggcg gcctgcgcgg ccttcaactt ctccaccagg tcatcaccca 9120 9180 gegeegege gatttgtace gggeeggatg gtttgegace gteaegeega tteeteggge ttgggggttc cagtgccatt gcagggccgg cagacaaccc agccgcttac gcctggccaa 9240 9300 ccgcccgttc ctccacacat ggggcattcc acggcgtcgg tgcctggttg ttcttgattt tocatgoogo otootttago ogotaaaatt catotactca tttattcatt tgotoattta 9360 9420 ctctggtagc tgcgcgatgt attcagatag cagctcggta atggtcttgc cttggcgtac 9480 cgcgtacatc ttcagcttgg tgtgatcctc cgccggcaac tgaaagttga cccgcttcat 9540 ggctggcgtg tctgccaggc tggccaacgt tgcagccttg ctgctgcgtg cgctcggacg gccggcactt agcgtgtttg tgcttttgct cattttctct ttacctcatt aactcaaatg 9600 9660 agttttgatt taatttcagc ggccagcgcc tggacctcgc gggcagcgtc gccctcgggt 9720 tetgatteaa gaaeggttgt geeggeggeg geagtgeetg ggtageteae gegetgegtg 9780 atacgggact caagaatggg cagctegtac ceggecageg ceteggeaac etcacegeeg

25136771 - 33 -

atgcgcgtgc	ctttgatcgc	ccgcgacacg	acaaaggccg	cttgtagcct	tccatccgtg	9840
acctcaatgc	gctgcttaac	cagctccacc	aggtcggcgg	tggcccatat	gtcgtaaggg	9900
cttggctgca	ccggaatcag	cacgaagtcg	gctgccttga	tcgcggacac	agccaagtcc	9960
gccgcctggg	gcgctccgtc	gatcactacg	aagtcgcgcc	ggccgatggc	cttcacgtcg	10020
cggtcaatcg	tcgggcggtc	gatgccgaca	acggttagcg	gttgatcttc	ccgcacggcc	10080
gcccaatcgc	gggcactgcc	ctggggatcg	gaatcgacta	acagaacatc	ggccccggcg	10140
agttgcaggg	cgcgggctag	atgggttgcg	atggtcgtct	tgcctgaccc	gcctttctgg	10200
ttaagtacag	cgataacctt	catgcgttcc	ccttgcgtat	ttgtttattt	actcatcgca	10260
tcatatacgc	agcgaccgca	tgacgcaagc	tgttttactc	aaatacacat	caccttttta	10320
gacggcggcg	ctcggtttct	tcagcggcca	agctggccgg	ccaggccgcc	agcttggcat	10380
cagacaaacc	ggccaggatt	tcatgcagcc	gcacggttga	gacgtgcgcg	ggcggctcga	10440
acacgtaccc	ggccgcgatc	atctccgcct	cgatctcttc	ggtaatgaaa	aacggttcgt	10500
cctggccgtc	ctggtgcggt	ttcatgcttg	ttcctcttgg	cgttcattct	cggcggccgc	10560
cagggcgtcg	gcctcggtca	atgcgtcctc	acggaaggca	ccgcgccgcc	tggcctcggt	10620
gggcgtcact	tcctcgctgc	gctcaagtgc	gcggtacagg	gtcgagcgat	gcacgccaag	10680
cagtgcagcc	gcctctttca	cggtgcggcc	ttcctggtcg	atcagctcgc	gggcgtgcgc	10740
gatctgtgcc	ggggtgaggg	tagggcgggg	gccaaacttc	acgcctcggg	ccttggcggc	10800
ctcgcgcccg	ctccgggtgc	ggtcgatgat	tagggaacgc	tcgaactcgg	caatgccggc	10860
gaacacggtc	aacaccatgc	ggccggccgg	cgtggtggtg	tcggcccacg	gctctgccag	10920
gctacgcagg	cccgcgccgg	cctcctggat	gcgctcggca	atgtccagta	ggtcgcgggt	10980
gctgcgggcc	aggcggtcta	gcctggtcac	tgtcacaacg	tcgccagggc	gtaggtggtc	11040
aagcatcctg	gccagctccg	ggcggtcgcg	cctggtgccg	gtgatcttct	cggaaaacag	11100
cttggtgcag	ccggccgcgt	gcagttcggc	ccgttggttg	gtcaagtcct	ggtcgtcggt	11160
gctgacgcgg	gcatagccca	gcaggccagc	ggcggcgctc	ttgttcatgg	cgtaatgtct	11220
ccggttctag	tcgcaagtat	tctactttat	gcgactaaaa	cacgcgacaa	gaaaacgcca	11280
ggaaaagggc	agggcggcag	cctgtcgcgt	aacttaggac	ttgtgcgaca	tgtcgttttc	11340
agaagacggc	tgcactgaac	gtcagaagcc	gactgcacta	tagcagcgga	ggggttggat	11400
caaagtactt	tgatcccgag	gggaaccctg	tggttggcat	gcacatacaa	atggacgaac	11460
ggataaacct	tttcacgccc	ttttaaatat	ccgttattct	aataaacgct	cttttctctt	11520
ag						11522

25136771 - 34 -

<210> 20 <211> 6610 <212> DNA <213> Artificial Sequence <220> <223> Vector pCI-n GL1-HEGO <400> 20 tcaatattgg ccattagcca tattattcat tggttatata gcataaatca atattggcta 60 ttggccattg catacgttgt atctatatca taatatgtac atttatattg gctcatgtcc 120 aatatgaccg ccatgttggc attgattatt gactagttat taatagtaat caattacggg 180 gtcattagtt catagcccat atatggagtt ccgcgttaca taacttacgg taaatggccc 240 gcctggctga ccgcccaacg acccccgccc attgacgtca ataatgacgt atgttcccat 300 agtaacgcca atagggactt tccattgacg tcaatgggtg gagtatttac ggtaaactgc 360 ccacttggca gtacatcaag tgtatcatat gccaagtccg ccccctattg acgtcaatga 420 cggtaaatgg cccgcctggc attatgccca gtacatgacc ttacgggact ttcctacttg 480 gcagtacatc tacgtattag tcatcgctat taccatggtg atgcggtttt ggcagtacac 540 caatgggcgt ggatagcggt ttgactcacg gggatttcca agtctccacc ccattgacgt 600 caatgggagt ttgttttggc accaaaatca acgggacttt ccaaaatgtc gtaataaccc 660 cgccccgttg acgcaaatgg gcggtaggcg tgtacggtgg gaggtctata taagcagagc 720 tegtttagtg aacegteaga teactagaag etttattgeg gtagtttate acagttaaat 780 tgctaacgca gtcagtgctt ctgacacaac agtctcgaac ttaagctgca gaagttggtc 840 gtgaggcact gggcaggtaa gtatcaaggt tacaagacag gtttaaggag accaatagaa 900 actgggcttg tcgagacaga gaagactctt gcgtttctga taggcaccta ttggtcttac 960 tgacatccac tttgcctttc tctccacagg tgtccactcc cagttcaatt acagctctta 1020 aggctagagt acttaatacg actcactata ggctagcgaa ggagatccgc catggcccac 1080 catcaccacc atcacggata tccatacgac gtgccagatt acgctcagtc gagtgccatg 1140 agcaagggcg aggaactgtt cactggcgtg gtcccaattc tcgtggaact ggatggcgat 1200 gtgaatgggc acaaattttc tgtcagcgga gagggtgaag gtgatgccac atacggaaag 1260 ctcaccctga aattcatctg caccactgga aagctccctg tgccatggcc aacactggtc 1320 actacettea cetatggegt geagtgettt tecagatace cagaceatat gaageageat 1380 gactttttca agagcgccat gcccgagggc tatgtgcagg agagaaccat ctttttcaaa 1440 gatgacggga actacaagac ccgcgctgaa gtcaagttcg aaggtgacac cctggtgaat 1500 agaatcgagc tgaagggcat tgactttaag gaggatggaa acattctcgg ccacaagctg 1560

25136771

1620 gaatacaact ataactccca caatgtgtac atcatggccg acaagcaaaa gaatggcatc 1680 aaggtcaact tcaagatcag acacaacatt gaggatggat ccgtgcagct ggccgaccat 1740 tatcaacaga acactccaat cggcgacggc cctgtgctcc tcccagacaa ccattacctg 1800 tccacccagt ctgccctgtc taaagatccc aacgaaaaga gagaccacat ggtcctgctg gagtttgtga ccgctgctgg gatcacacat ggcatggacg agctgtacaa gggcgccggc 1860 1920 gctggtgctg gtgctggcgc catcagcgcg ttgaccctcc acaccaaagc atctgggatg 1980 gccctactgc atcagatcca agggaacgag ctggagcccc tgaaccgtcc gcagctcaag atccccctgg agcggcccct gggcgaggtg tacctggaca gcagcaagcc cgccgtgtac 2040 2100 aactaccccg agggcgccgc ctacgagttc aacgccgcgg ccgccgccaa cgcgcaggtc tacggtcaga ccggcctccc ctacggcccc gggtctgagg ctgcggcgtt cggctccaac 2160 2220 ggcctggggg gtttcccccc actcaacagc gtgtctccga gcccgctgat gctactgcac 2280 ccgccgccgc agctgtcgcc tttcctgcag ccccacggcc agcaggtgcc ctactacctg gagaacgagc ccagcggcta cacggtgcgc gaggccggcc cgccggcatt ctacaggcca 2340 2400 aattcagata atcgacgcca gggtggcaga gaaagattgg ccagtaccaa tgacaaggga 2460 agtatggcta tggaatctgc caaggagact cgctactgtg cagtgtgcaa tgactatgct tcaggctacc attatggagt ctggtcctgt gagggctgca aggccttctt caagagaagt 2520 attcaaggac ataacgacta tatgtgtcca gccaccaacc agtgcaccat tgataaaaac 2580 2640 aggaggaaga gctgccaggc ctgccggctc cgcaaatgct acgaagtggg aatgatgaaa 2700 ggtgggatac gaaaagaccg aagaggaggg agaatgttga aacacaagcg ccagagagat 2760 gatggggagg gcaggggtga agtggggtct gctggagaca tgagagctgc caacctttgg 2820 ccaagcccgc tcatgatcaa acgctctaag aagaacagcc tggccttgtc cctgacggcc 2880 gaccagatgg tcagtgcctt gttggatgct gagcccccca tactctattc cgagtatgat 2940 cctaccagac ccttcagtga agcttcgatg atgggcttac tgaccaacct ggcagacagg 3000 gagctggttc acatgatcaa ctgggcgaag agggtgccag gctttgtgga tttgaccctc 3060 catgatcagg tccaccttct agaatgtgcc tggctagaga tcctgatgat tggtctcgtc tggcgctcca tggagcaccc agtgaagcta ctgtttgctc ctaacttgct cttggacagg 3120 3180 aaccagggaa aatgtgtaga gggcatggtg gagatcttcg acatgctgct ggctacatca tctcggttcc gcatgatgaa tctgcaggga gaggagtttg tgtgcctcaa atctattatt 3240 ttgcttaatt ctggagtgta cacatttctg tccagcaccc tgaagtctct ggaagagaag 3300 3360 gaccatatcc accgagtcct ggacaagatc acagacactt tgatccacct gatggccaag 3420 gcaggcctga ccctgcagca gcagcaccag cggctggccc agctcctcct catcctctcc

25136771 - 36 -

(	cacatcaggc	acatgagtaa	caaaggcatġ	gagcatctgt	acagcatgaa	gtgcaagaac	3480
9	gtggtgcccc	tctatgacct	gctgctggag	atgctggacg	cccaccgcct	acatgcgccc	3540
ě	actagccgtg	gaggggcatc	cgtggaggag	acggaccaaa	gccacttggc	cactgcgggc	3600
1	tctacttcat	cgcattcctt	gcaaaagtat	tacatcacgg	gggaggcaga	gggtttccct	3660
•	gccacagtct	gagagetece	tggcggaatt	cggatcgggc	ggccgcttcg	agcagacatg	3720
ě	ataagataca	ttgatgagtt	tggacaaacc	acaactagaa	tgcagtgaaa	aaaatgcttt	3780
ě	atttgtgaaa	tttgtgatgc	tattgcttta	tttgtaacca	ttataagctg	caataaacaa	3840
•	gttaacaaca	acaattgcat	tcattttatg	tttcaggttc	agggggagat	gtgggaggtt	3900
	tttaaagca	agtaaaacct	ctacaaatgt	ggtaaaatcg	ataaggatcc	gggctggcgt	3960
•	aatagcgaag	aggcccgcac	cgatcgccct	tcccaacagt	tgcgcagcct	gaatggcgaa	4020
	tggacgcgcc	ctgtagcggc	gcattaagcg	cggcgggtgt	ggtggttacg	cgcagcgtga	4080
,	ccgctacact	tgccagcgcc	ctagcgcccg	ctcctttcgc	tttcttccct	tcctttctcg	4140
	ccacgttcgc	cggctttccc	cgtcaagctc	taaatcgggg	gctcccttta	gggttccgat	4200
	ttagagcttt	acggcacctc	gaccgcaaaa	aacttgattt	gggtgatggt	tcacgtagtg	4260
,	ggccatcgcc	ctgatagacg	gtttttcgcc	ctttgacgtt	ggagtccacg	ttctttaata	4320
	gtggactctt	gttccaaact	ggaacaacac	tcaaccctat	ctcggtctat	tcttttgatt	4380
	tataagggat	tttgccgatt	tcggcctatt	ggttaaaaaa	tgagctgatt	taacaaatat	4440
	ttaacgcgaa	ttttaacaaa	atattaacgt	ttacaatttc	gcctgatgcg	gtattttctc	4500
	cttacgcatc	tgtgcggtat	ttcacaccgc	atatggtgca	ctctcagtac	aatctgctct	4560
	gatgccgcat	agttaagcca	gccccgacac	ccgccaacac	ccgctgacgc	gccctgacgg	4620
	gcttgtctgc	tcccggcatc	cgcttacaga	caagctgtga	ccgtctccgg	gagctgcatg	4680
	tgtcagaggt	tttcaccgtc	atcaccgaaa	cgcgcgagac	gaaagggcct	cgtgatacgc	4740
	ctatttttat	aggttaatgt	catgataata	atggtttctt	agacgtcagg	tggcactttt	4800
	cggggaaatg	tgcgcggaac	ccctatttgt	ttatttttct	aaatacattc	aaatatgtat	4860
	ccgctcatga	gacaataacc	ctgataaatg	cttcaataat	attgaaaaag	gaagagtatg	4920
	agtattcaac	atttccgtgt	cgcccttatt	cccttttttg	cggcattttg	ccttcctgtt	4980
	tttgctcacc	cagaaacgct	ggtgaaagta	aaagatgctg	aagatcagtt	gggtgcacga	5040
	gtgggttaca	tcgaactgga	tctcaacagc	ggtaagatcc	ttgagagttt	tcgccccgaa	5100
	gaacgttttc	caatgatgag	cacttttaaa	gttctgctat	gtggcgcggt	attatcccgt	5160
	attgacgccg	ggcaagagca	actcggtcgc	cgcatacact	attctcagaa	tgacttggtt	5220

25136771 - 37 -

gagtactcac	cagtcacaga	aaagcatctt	acggatggca	tgacagtaag	agaattatgc	5280
agtgctgcca	taaccatgag	tgataacact	gcggccaact	tacttctgac	aacgatcgga	5340
ggaccgaagg	agctaaccgc	ttttttgcac	aacatggggg	atcatgtaac	tcgccttgat	5400
cgttgggaac	cggagctgaa	tgaagccata	ccaaacgacg	agcgtgacac	cacgatgcct	5460
gtagcaatgg	caacaacgtt	gcgcaaacta	ttaactggcg	aactacttac	tctagcttcc	5520
cggcaacaat	taatagactg	gatggaggcg	gataaagttg	caggaccact	tctgcgctcg	5580
gcccttccgg	ctggctggtt	tattgctgat	aaatctggag	ccggtgagcg	tgggtctcgc	5640
ggtatcattg	cagcactggg	gccagatggt	aagccctccc	gtatcgtagt	tatctacacg	5700
acggggagtc	aggcaactat	ggatgaacga	aatagacaga	tcgctgagat	aggtgcctca	5760
ctgattaagc	attggtaact	gtcagaccaa	gtttactcat	atatacttta	gattgattta	5820
aaacttcatt	tttaatttaa	aaggatctag	gtgaagatcc	tttttgataa	tctcatgacc	5880
aaaatccctt	aacgtgagtt	ttcgttccac	tgagcgtcag	accccgtaga	aaagatcaaa	5940
ggatcttctt	gagatccttt	ttttctgcgc	gtaatctgct	gcttgcaaac	aaaaaacca	6000
ccgctaccag	cggtggtttg	tttgccggat	caagagctac	caactctttt	tccgaaggta	6060
actggcttca	gcagagcgca	gataccaaat	actgtccttc	tagtgtagcc	gtagttaggc	6120
caccacttca	agaactctgt	agcaccgcct	acatacctcg	ctctgctaat	cctgttacca	6180
gtggctgctg	ccagtggcga	taagtcgtgt	cttaccgggt	tggactcaag	acgatagtta	6240
ccggataagg	cgcagcggtc	gggctgaacg	gggggttcgt	gcacacagcc	cagcttggag	6300
cgaacgacct	acaccgaact	gagataccta	cagcgtgagc	tatgagaaag	cgccacgctt	6360
cccgaaggga	gaaaggcgga	caggtatccg	gtaagcggca	gggtcggaac	aggagagcgc	6420
acgagggagc	ttccaggggg	aaacgcctgg	tatctttata	gtcctgtcgg	gtttcgccac	6480
ctctgacttg	agcgtcgatt	tttgtgatgc	tcgtcagggg	ggcggagcct	atggaaaaac	6540
gccagcaacg	cggccttttt	acggttcctg	gccttttgct	ggccttttgc	tcacatggct	6600
cgacagatct						6610

<210> 21

<211> 2525

<212> DNA

<213> Violet Cress

<400> 21

agcaatggtt caacatcaac caccacca agtccctct cctccgtcgc agcaatctcc 60 ggtaacacca ctaacggcgg cgtttgggat gagattaggt ggtctagaag gtttgttcgg 120 tccttacggg atacgtttt acacggcggc gaagatagcc gagttaggtt tcacggcgag 180

25136771 - 38 -

240 cactctcgtt ggtatgaaag acgaagagct tgaagatatg atgaatagtc tctctcatat ctttcgttgg gagcttcttg ttggtgaacg ttacggtata aaagctgccg ttagagctga 300 acggaggaga ttgcaagaag aggaggaaga tgaatcttct agacgccgtc atttgcttct 360 ctccgccgct ggtgattccg gcactcacca cgctcttgat gctctctctc aagaaggtac 420 taatattaaa atttatataa ttttttttca cggtttatat atttacgtac ttattgttat 480 acgtatcacg taagtaactt atctatatag gttcaataaa ctgttaaaaa tctatgttca 540 cgtgaccgat tgatataaag ttatatgagt ttttacatga tatatattag agtaatattg 600 tatgatacct agtcgttaac aacataattt aatgaatcat tataaaataa tattgcaatt 660 gttaaggaat ctctattcac attcaacact tctatataaa ttaacttaaa aattgttgac 720 aaacaaaaaa ttaacttaaa aataataaac tcattatgta tatagtatgt gtatagtaac 780 gctattatct atagagtttc acgagtgata gtctgaattt tatttattaa agtcctatta 840 caaagtacga ggaataatat catataatat aattatatac gcgtatatat atttaaattt 900 ttaatgaacg tgacataagt aactgtagat gattggacag gcttatctga ggaaccgatg 960 catcaagacc aaactgacgc ggcgggtaac ggcggattcg gtggttattt ggaatcatca 1020 gtacacggaa agatgaagaa acatcaacca agacgtagaa agaaaccgtt ggtactgacg 1080 tcagttgaaa ccgacgatga cggcaacgat aacgaggatg acgacgggat ggataacggt 1140 aacggaggta ttgggttagg gacggagaga cagagagaac atccgtttat tgtaactgag 1200 cctggggaag tggcacgtgg caaaaagaac ggtttggatt atcttttcca cttgtacgaa 1260 caatgccgtg agttccttct tcaggtccag actattgcta aagaccgtgg cgaaaaatgc 1320 cccaccaagg tatctttctc aagatttaat actagccgtt gatttaattc aattaaatat 1380 1440 taatcatgga cagctgtttt aaagtaagtt tgacaagtca cgagttcaga gttttcacta 1500 ggattttttg tttagttatt gctatgtctg gcaaacattt atataggttt tagatttgtt 1560 taattttttt ttattctacc tttagaataa atatagagtt agattttcaa tataaaaaat 1620 gtgacagtca ccgtacaact acgtgtcgag ttagggaaag gagacagttg tttcttgttt 1680 ttggctaaaa cgacataaga tagcaagata cgatgtcctc atgctttaga tactatttct 1740 gcatgcattc acacacacat gtataatgtt taggtgtata tattagctag atatttctat 1800 ttttataatt tgttgggtgc atatatatat taaaaaatag tatagtatat agtgataata 1860 tttagaaaca gtctattgat cagcgggtaa taaagggtct ctactttcca ctagtgaaga 1920 tagtacaaaa tccacgagct ctgtgccatt gcttgtacag acgcacgtta ctaaattctg 1980 agttgttaca gaaaatttaa ccgaataaaa ctctttttaa tttcttcttt caaaaatttg 2040

25136771 - 39 -

ggttttgaa	c atatttttt	tggctgttta	gtctaacgaa	gagetgtatt	aaagatactg	2100
tatttatca	c tactaaattt	attttgaaaa	tctagtataa	ataaattatt	tacttcatat	2160
tgattatta	t ggtaatgaaa	ggttatatgt	aaaacaggtg	acgaaccaag	tgtttaggta	2220
cgctaaaaa	a tcaggagcga	gttacataaa	caaaccaaaa	atgcgacact	acgttcattg	2280
ttacgctct	c cactgcctag	acgaagaagc	atcaaacgct	ctaagaagag	cgtttaaaga	2340
acgcggcga	g aacgttggct	cgtggcgtca	ggcttgttac	aagccgctag	tgaacatagc	2400
ctgtcgtca	t ggctgggaca	tagacgccgt	tttcaacgca	catcctcgtc	tatctatttg	2460
gtacgttcc	a actaaactgc	gtcagctttg	ccatttggag	cgtaacaacg	ccgttgctgc	2520
ggcgg						2525
<210> 22 <211> 9 <212> DN <213> Hu <400> 22 ggggcgtgg	A man					9
ggggcgcgg						9
<212> DN	78					
<400> 23		**			<b>.</b>	60
	a ttcgagcaca					120
	g aacgtttctg t gcgtgcgcgg					120 180
	c cacaccaaag					240
	c ctcagccgcc					300
	c agcagtaagc					360
	c accgccccg					420
	g accgccttcg					480
	c ccgctgatgc					540
	c cagcaggtgc					600
	c cctccagcct					660
	g tccagcagca					720
	t gcagtgtgca					780

25136771 - 40 -

tgaa	ggctgc	aaggctttct	tcaagagaag	tattcaagga	cacaatgact	acatgtgtcc	840
agct	acaaac	caatgcacaa	tcgacaagaa	caggagaaag	agctgccagg	cctgcaggct	900
gcgc	aagtgt	tacgaagtag	gcatgatgaa	aggtgggata	cggaaagacc	ggagaggagg	960
gaga	atgctg	aaacacaagc	gccagagaga	cgacttggaa	ggcaggaacg	acatggggcc	1020
ttca	ggagac	atgagggcca	ccaacctttg	gcctagtcct	cttgtgatta	agcacactaa	1080
gaag	aacagc	cctgccttgt	ccttgacagc	cgaccaaatg	gtcagtgcct	tgttggatgc	1140
tgaa	ccgccc	ttaatctatt	ctgaatatga	tccttctaga	cctttcagcg	aagcttcgat	1200
gato	ggatta	ttgaccaacc	tggcagacag	ggagttggtt	catatgatca	actgggcaaa	1260
gaga	igtgcca	ggctttggag	acttaaatct	ccatgatcag	gtccacctcc	tggagtgtgc	1320
ctgç	ıttggag	atcctgatga	ttggtctcat	ctggcgctcc	atggaacacc	cagggaagct	1380
cctç	rtttgct	cctaatttgc	tcctggacag	gaatcagggc	aagtgtgtgg	agggcatggt	1440
ggag	atcttt	gacatgttgc	tggctacatc	agctcggttc	cgcatgatgg	acctgcaggg	1500
agag	gagttt	gtgtgcctca	aatctatcat	tttgcttaat	tctggagtgt	acacatttct	1560
gtco	agcacc	ttgaagtctc	tggaggagaa	ggaccacatc	caccgggtcc	tggataagat	1620
caca	igacact	ttgattcacc	tgatggccaa	agctggcctg	acactgcagc	agcagcatcg	1680
tcgt	ctggcc	cagctcctcc	tcattctttc	ccacatccgg	cacatgagta	acaaaggcat	1740
ggag	cacctc	tacaacatga	aatgcaagaa	cgttgtaccc	ttctatgacc	tgttgctgga	1800
gato	ıttggat	gctcaccgcc	tgcatacccc	cgtcagtcgc	atgggggtct	ccccagagga	1860
gccc	agtcag	agccagctga	ccaccaccaa	ctccacttca	tcacattcct	tacaaaccta	1920
ctac	atcccg	tcggaagcag	agagtttccc	caacacaatc	tgagacctcc	caggctcc	1978

<210> 24 <211> 1788

<212> DNA

<213> Pig

<400> 24

atgaccatga coctacacac caaagcgtcc ggcatggccc tgctgcacca gatccaagcc 60
aacgagctgg agcccctgaa ccgcccgcag ctcaagatcc ccctggagcg gcccctgggc 120
gaggtgtacg tggacagcag caagcccgcc gtgtataact accccgaggg cgccgcgtac 180
gacttcaacg ccgcggccgc cgcctccgcg cccgtctacg gccagtcggg cctcgcctac 240
ggcccggggt cggaggcggc ggcgttcggc gccaacggct tggggggctt ccagccgct 300
aacagcgtgt ctccgagccc gctggtgctg ctgcacccgc cgccgcagct ctcgccttc 360
ctgcaccccc acggccaaca ggtgccctat tacctggaga atgagccgag cggctatgcg 420

25136771 - 41 -

gtgcgcgagg	ccggccctcc	cgccttctac	aggccaaatt	cagataatcg	gcgccagggt	480
ggcagagaga	gattggccag	caccagtgac	aagggaagca	tggccatgga	atctgccaag	540
gagactcgct	actgtgcagt	gtgcaatgac	tatgcctcag	gctaccatta	tggagtttgg	600
tcttgcgagg	gctgtaaggc	cttcttcaag	agaagtattc	aaggacataa	tgactacatg	660
tgtccagcta	ccaaccagtg	cacaattgat	aagaacagga	ggaagagctg	tcaggcctgc	720
cggctacgca	agtgctacga	agtgggcatg	atgaaagggg	ggatacggaa	agaccggaga	780
ggagggagaa	tgttgaagca	caagcgccag	agagatgatg	gagagggcag	gaatgaagcg	840
gtgccccctg	gagacatgag	atctgccaac	ctttggccaa	gccctctctt	gattaaacac	900
actaagaaga	acagcccggt	cttgtccctg	acagccgacc	agatgatcag	tgccttgttg	960
gaggctgagc	cccccataat	ctattccgag	tatgatccta	ccagacccct	cagtgaggct	1020
tcaatgatgg	gcttgctgac	caacctcgca	gacagggagc	tggtacacat	gatcaactgg	1080
gcaaagaggg	tgccaggatt	tttggattta	agcctccatg	atcaagtgca	tcttctggaa	1140
tgtgcctggc	tagagatcct	catgattggt	cttgtctggc	gctccatgga	gcacccaggg	1200
aagctcctgt	ttgctcctaa	cttgctcctg	gacaggaacc	agggcaagtg	tgtcgaggga	1260
atggtggaga	tctttgacat	gttgctggct	acatcatctc	gcttccgtat	gatgaatctc	1320
cagggagagg	agtttgtgtg	cctcaaatcc	atcattttgc	ttaattctgg	agtgtacacg	1380
tttctgtcca	gcaccctgaa	gtctctggaa	gagaaggacc	atatccaccg	tgtcctggac	1440
aagatcacag	acaccttgat	ccacctgatg	gccaaagcgg	gcctgactct	gcagcagcag	1500
caccggcgtc	tcgcgcagct	cctcctcatc	ctgtctcact	tcaggcacat	gagtaacaaa	1560
ggcatggagc	atctgtacaa	catgaagtgc	aagaacgtgg	tgcccctcta	tgacctgctg	1620
ctggagatgc	tggacgccca	ccgcctgcac	gccccaacca	accttggggg	cccacccccg	1680
gaggacatga	gccagagcca	gctggccacc	tcgggctcaa	ctccatcgca	ttccttgcaa	1740
atgtattaca	tcacagggga	ggcggagaac	ttccccacca	caatctga		1788

<210> 25

<211> 1759

<212> DNA

<213> Polyxenus fasciculatus

<400> 25

tcatatgctt gtctcaaaga ttaagccatg catgtgtaag tacacaccat cctaaggtga 60 gaccgcgaat ggctcattaa atcagttatg gttcattaga tgagtccaat cctacttgga 120 taactgtggc aattctagag ctaatacacg cctccaagct ctgacctatc gggacgagcg 180 cttttattag accaagacca atcgggcttc ggtccgtttc ctttggtgac tctgaataac 240

25136771 - 42 -

cttttgaaga	tcgcacggtc	tcgaaccggc	gatgcatctt	tcaaatgtct	gccttatcaa	300
ctgtcgatgg	taagttatgc	gcttaccatg	gttgtaacgg	gtaacggaga	atcagggttc	360
gattccggag	agggagcctg	agaaacggct	gccacatcca	aggaaggcag	caggcacgca	420
aattacccac	tcccggcacg	gggaggtagt	gacgaaaaat	aacgatgcgg	gactcttccg	480
aggccccgta	atcggaatga	gtacacttta	aatcctttaa	cgaggatcaa	ttggagggca	540
agtctggtgc	cagcagccgc	ggtaattcca	gctccaatag	cgtatactaa	agttgttgcg	600
gctaaaaagc	tcgtagttgg	atttcagtcg	taggccggtg	gtccaccgcc	cggtggctac	660
tgcctggtct	ggacaccttg	ccagctctcc	ggcgatgctc	ttgaccgggt	gtcgttggtg	720
gctggaacgt	ttactttgaa	aaaattagag	tgctctaagc	aggtgctatc	ggcttgaata	780
acacagcatg	gaataatgga	acacgacctt	ggttctgttc	tgttggtctt	tggaagccaa	840
ggtaatgatt	aatagggacg	gacgggggca	ttcgtattgc	gacgctagag	gtgaaattct	900
tggaccgtcg	caagacgaac	tactgcgaac	gcatttgcca	agaacgtttt	cattaatcaa	960
gaacgaaagt	cagaggttcg	aaggcgatca	gataccgccc	tagttctgac	cataaacaat	1020
gccaaccagc	gatccgccgg	agttactccc	atgactcggc	gggcagcttc	cgggaaacca	1080
aagtgtttgg	gttccggggg	aagtatggtt	gcaaagctga	aacttaaagg	aattgacgga	1140
agggcaccac	caggagtgga	gcctgcggct	taatttgact	caacacggga	cacctcaccc	1200
ggcccggaca	ccggaaggat	tgacagactg	agagctcttt	cttgattcgg	tgggtggtgg	1260
tgcatggccg	ttcttagttg	gtggagcgat	ctgtctggtt	aattccgata	acgaacgaga	1320
ctctagccta	ctaactagcc	agtcgatcat	ttgtcggctg	ttcttcttag	agggataagc	1380
ggcttttagc	cgcatgagat	tgagcaataa	caggtctgtg	atgcccttag	atgtccggga	1440
ccgcacgcgc	gctacactga	tgttgtcagc	ttgtttctcc	ccttgtccga	gaggaccggg	1500
taatccgctg	aaccaccttc	gtgataggga	tcggggtttg	aaattatccc	ccgtgaacga	1560
ggaattccca	gtaagcgcga	gtcataagct	cgtgttgatt	acgtccctgc	cctttgtaca	1620
caccgcccgt	cgctactacc	gattgaatga	tttagtgagg	tcttcggact	gaggcccggc	1680
gaagcttgct	ttgccgacgc	tttggaaaga	tgatcgaact	tgatcattta	gaggaagtaa	1740
aagtcgtaac	aaggtttcc					1759

<210> 26 <211> 7 <212> DNA <213> Human

<400> 26 ccgcccc

25136771 - 43 -

<210> <211> <212> <213>	27 228 DNA Poly	yxenus lagur	cus				
<400> tcatttq	27 gtcg	gctgttcttc	ttagagggat	aagcggcttt	tagccgcatg	agattgagca	60
ataacag	ggtc	tgtgatgccc	ttagatgtcc	gggaccgcac	gcgcgctaca	ctgatgttat	120
cagctto	gttt	ctccccttgt	ccgagaggac	cgggtaatcc	gctgaaccac	cttcgtgata	180
gggatco	aggg	tttgaaatta	tcccccgtga	acgaggaatt	cccagtaa		228
<210> <211> <212> <213>	28 2842 DNA Huma						
<400> acacgca	28 acct	catcgacctg	gtaaccaatg	agattatcac	ccaaggacct	ccagtggact	60
ggaatga	acat	tgctggtctc	gacctggtga	aggctgtcat	taaagaggag	gttttatggc	120
cagtgtt	gag	gtcagacgcg	ttcagtggac	tgacggcctt	acctcggagc	atccttttat	180
ttggaco	ctcg	ggggacaggc	aaaacattat	tgggcagatg	catcgctagt	cagctggggg	240
ccacatt	ttt	caaaattgcc	ggttctggac	tagtcgccaa	gtggttagga	gaagcagaga	300
aaattat	cca	tgcctctttt	cttgtggcca	ggtgtcgcca	gccctcggtg	atttttgtta	360
gtgacat	tga	catgcttctc	tcctctcaag	tgaatgagga	acatagtcca	gtcagtcgga	420
tgagaad	ccga	atttctgatg	caactggaca	ctgtactaac	ttcggctgag	gaccaaatcg	480
tagtaat	ttg	tgccaccagt	aaaccagaag	aaatagatga	atcccttcgg	aggtacttca	540
tgaaac	gact	tttaatccca	cttcctgaca	gcacagcgag	gcaccagata	atagtacaac	600
tgctctc	caca	gcacaattac	tgtctcaatg	acaaggagtt	tgcactgctc	gtccagcgca	660
cagaag	gctt	ttctggacta	gatgtggctc	atttgtgtca	ggaagcagtg	gtgggccccc	720
tccatgo	ccat	gccagccaca	gacctttcag	ccattatgcc	cagccagttg	aggcccgtta	780
catatca	aaga	ctttgaaaat	gctttctgca	agattcagcc	tagcatatct	caaaaggagc	840
ttgatat	tgta	tgttgaatgg	aacaaaatgt	ttggttgcag	tcagtgataa	cttctttaga	900
aaaaaa	aaat	gtaatgaatg	ttggcacaca	cacataaaac	ctgctacata	gggaatagag	960
ccccttt	cca	gtagagttta	aattgcaaag	ggtactgggg	aagatgacga	ttaagttgca	1020
tctttag	gagt	cagggtagat	ttggaggaaa	agtgcatcaa	atgagagctt	ctgatttgaa	1080
agcccca	agat	gacagaaagc	atatgtggat	gctcagttct	gttcaagcta	gacaacactc	1140
accaago	gagc	aaggtgcaag	tgtgttgatt	tcagaaggac	atgaacctcg	tgtgttgatt	1200

25136771 - 44 -

ccattctgct	gttctcgaga	tttagttgct	gtcaagtgcc	tggagtggtg	ctttattttt	1260
tgtttgcctc	acaattacat	tggtggcatg	tgctaatata	aagagcttta	acttcaaaca	1320
ttattggact	aaagagatga	acagttgtgt	tatgacagaa	aaccagattt	ttgccatttt	1380
aagagcaaca	gtattcctca	atcctgtctg	ttctgcagta	ttaagctaag	aacaggtaaa	1440
acagggtaac	ggtaatctgg	accttaattt	ctgcagttca	tttcttttaa	tgttcttgtc	1500
tgcaaaaact	caggaaagtg	attgtgattt	gtacagtacc	tcaaaggaat	gtgttgaaag	1560
cactatgtac	tgctgagagt	aataggatag	gcttcaatgt	tactttatat	taaaatgtat	1620
gtttacctca	acaattggaa	aatagcaagg	aaaattactt	tgaatgtatc	cagaaaaata	1680
ctgaagtgtg	atacaactga	atatttacag	tttaaagtag	aaatggaagg	atttttttaa	1740
gttcttttac	taattatggg	gaattaacca	gagcagaata	attctttatg	tcaataactg	1800
caagagttct	tagtacattg	ctccttgata	attaagtgaa	aatgttctta	aaaggtacac	1860
tggttaattg	aaagctactt	attcagtttg	tgttagtgtc	tagacctgtc	agccacaaga	1920
cctgtttagg	accctgaaag	tcacagtacc	taaaaactat	gactgccttt	ttattgcata	1980
ggtggtagtg	gtggtgatgg	tggtggtagt	ttgcaagtta	tctcttaaaa	ctgctgggaa	2040
tggtgtcatt	ctattcacta	atctagctta	tagacttgcc	gtgctgtttg	atagaatgca	2100
gaggatagca	accaaaacaa	atacacaaat	aaataaaaac	aaaaaccaac	caacaaacca	2160
acttacatac	atatatatat	atccacaaag	aacctctcca	tctcctcccc	ttctttttga	2220
ctccactctt	gtcagtgcaa	ttttgcttct	cattttgaaa	tctgggctgt	agtgctcctg	2280
ctttatttct	acctcagttt	tgttacattt	ctcttggaaa	gtaaagtaga	aaattggaag	2340
tggacacaca	cactgcaatg	tagcttgcca	aacatgttac	tttgttttct	tccatctttc	2400
accgtaaatc	tagtttccaa	agacatcagc	atttgtgctt	acttccacct	cagtctacca	2460
gccccacccc	tacccatggc	ataagtggca	tttttcttaa	tttcctattt	ttctcctgct	2520
ctctgtcaag	ttgttctttg	tatcctttaa	tgctttatgt	gcaacctttc	attgatagtg	2580
ggctgatgtt	tggcaatgct	tctgaactgt	cacagagcag	gctgtagctt	tccacagcca	2640
ctgcccatgc	ataagcagaa	cagcctggcc	ttttgaatgt	attttcctgg	gtttttttcc	2700
ccttttcttt	ttttagttta	gagatgcagt	aacaaaactg	ttgcaaagca	ctggcatttt	2760
atgtattcaa	taaataagtg	atgtacattt	ttaaaaaaat	ttaaataaat	gcaatgagaa	2820
gccccaaaaa	aaaaaaaaa	aa				2842

25136771 - 45 -

<sup>&</sup>lt;210> 29 <211> 381 <212> PRT

<213> violet cress

<400> 29

Ala Met Val Gln His Gln Pro Pro Pro Gln Val Pro Pro Pro Pro Ser 1 5 10 15

Gln Gln Ser Pro Val Thr Pro Leu Thr Ala Ala Phe Gly Met Arg Leu 20 25 30

Gly Gly Leu Glu Gly Leu Phe Gly Pro Tyr Gly Ile Arg Phe Tyr Thr 35 40 45

Ala Ala Lys Ile Ala Glu Leu Gly Phe Thr Ala Ser Thr Leu Val Gly 50 55 60

Met Lys Asp Glu Glu Leu Glu Asp Met Met Asn Ser Leu Ser His Ile 65 70 75 80

Phe Arg Trp Glu Leu Leu Val Gly Glu Arg Tyr Gly Ile Lys Ala Ala 85 90 95

Val Arg Ala Glu Arg Arg Leu Gln Glu Glu Glu Glu Asp Glu Ser 100 105 110

Ser Arg Arg Arg His Leu Leu Ser Ala Ala Gly Asp Ser Gly Thr 115 120 125

His His Ala Leu Asp Ala Leu Ser Gln Glu Asp Asp Trp Thr Gly Leu 130 135 140

Ser Glu Glu Pro Met His Gln Asp Gln Thr Asp Ala Ala Gly Asn Gly 145 150 155 160

Gly Phe Gly Gly Tyr Leu Glu Ser Ser Val His Gly Lys Met Lys Lys 165 170 175

His Gln Pro Arg Arg Arg Lys Lys Pro Leu Val Leu Thr Ser Val Glu 180 185 190

Thr Asp Asp Asp Gly Asn Asp Asn Glu Asp Asp Asp Gly Met Asp Asn 195 200 205

Gly Asn Gly Gly Ile Gly Leu Gly Thr Glu Arg Gln Arg Glu His Pro 210 215 220

Phe Ile Val Thr Glu Pro Gly Glu Val Ala Arg Gly Lys Lys Asn Gly

25136771 - 46 -

225 230 235 240

Leu Asp Tyr Leu Phe His Leu Tyr Glu Gln Cys Arg Glu Phe Leu Leu 245 250 255

Gln Val Gln Thr Ile Ala Lys Asp Arg Gly Glu Lys Cys Pro Thr Lys 260 265 270

Val Thr Asn Gln Val Phe Arg Tyr Ala Lys Lys Ser Gly Ala Ser Tyr 275 280 285

Ile Asn Lys Pro Lys Met Arg His Tyr Val His Cys Tyr Ala Leu His 290 295 300

Cys Leu Asp Glu Glu Ala Ser Asn Ala Leu Arg Arg Ala Phe Lys Glu 305 310 315 320

Arg Gly Glu Asn Val Gly Ser Trp Arg Gln Ala Cys Tyr Lys Pro Leu 325 330 335

Val Asn Ile Ala Cys Arg His Gly Trp Asp Ile Asp Ala Val Phe Asn 340 345 350

Ala His Pro Arg Leu Ser Ile Trp Tyr Val Pro Thr Lys Leu Arg Gln 355 360 365

Leu Cys His Leu Glu Arg Asn Asn Ala Val Ala Ala Ala 370 375 380

<210> 30

<211> 595

<212> PRT

<213> hamster

<400> 30

Met Thr Met Thr Leu His Thr Lys Ala Ser Gly Met Ala Leu Leu His 1 5 10 15

Gln Ile Gln Gly Asn Glu Leu Glu Pro Leu Ser Arg Pro Gln Leu Lys 20 25 30

Met Pro Leu Glu Arg Ala Leu Ser Glu Val Tyr Val Asp Ser Ser Lys 35 40 45

Pro Ala Met Phe Asn Tyr Pro Glu Gly Ala Ala Tyr Glu Phe Asn Ala 50 55 60

25136771 - 47 -

Ala Thr Ala Pro Ala Pro Val Tyr Gly Gln Thr Gly Ile Ala Tyr Gly Ser Gly Ser Glu Ala Thr Ala Phe Gly Ser Asn Ser Leu Gly Leu Phe Pro Gln Leu Asn Ser Val Ser Pro Ser Pro Leu Met Leu Leu His Pro 100 105 Pro Pro Gln Leu Ser Pro Phe Leu His Pro His Gly Gln Gln Val 120 115 Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Ala Tyr Ala Val Arg Asp Ser 130 135 Gly Pro Pro Ala Phe Tyr Arg Ser Asn Ser Asp Asn Arg Arg Gln Ser 150 155 Gly Arg Glu Arg Leu Ser Ser Ser Glu Lys Gly Ser Met Ala Met Glu Ser Val Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala 185 Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe Phe.Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 235 Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg 245 Lys Asp Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 265 Asp Leu Glu Gly Arg Asn Asp Met Gly Pro Ser Gly Asp Met Arg Ala 280

Thr Asn Leu Trp Pro Ser Pro Leu Val Ile Lys His Thr Lys Lys Asn

295

25136771 - 48 -

Ser Pro Ala Leu Ser Leu Thr Ala Asp Gln Met Val Ser Ala Leu Leu 310 315 Asp Ala Glu Pro Pro Leu Ile Tyr Ser Glu Tyr Asp Pro Ser Arg Pro 325 330 Phe Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 345 340 Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Gly 355 360 Asp Leu Asn Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 370 375 Glu Ile Leu Met Ile Gly Leu Ile Trp Arg Ser Met Glu His Pro Gly 395 Lys Leu Leu Phe Ala Pro Asn Leu Leu Asp Arg Asn Gln Gly Lys 410 Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 425 Ala Arg Phe Arg Met Met Asp Leu Gln Gly Glu Glu Phe Val Cys Leu Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 485 Leu Gln Gln His Arg Arg Leu Ala Gln Leu Leu Ile Leu Ser 500 His Ile Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Asn Met Lys Cys Lys Asn Val Val Pro Phe Tyr Asp Leu Leu Glu Met Leu 530 535 Asp Ala His Arg Leu His Thr Pro Val Ser Arg Met Gly Val Ser Pro

25136771 - 49 -

545 550 555 560

Glu Glu Pro Ser Gln Ser Gln Leu Thr Thr Thr Asn Ser Thr Ser Ser 565 570 575

His Ser Leu Gln Thr Tyr Tyr Ile Pro Ser Glu Ala Glu Ser Phe Pro 580 585 590

Asn Thr Ile 595

<210> 31

<211> 595

<212> PRT

<213> pig

<400> 31

Gln Ile Gln Ala Asn Glu Leu Glu Pro Leu Asn Arg Pro Gln Leu Lys 20 25 30

Ile Pro Leu Glu Arg Pro Leu Gly Glu Val Tyr Val Asp Ser Ser Lys 35 40 45

Pro Ala Val Tyr Asn Tyr Pro Glu Gly Ala Ala Tyr Asp Phe Asn Ala 50 60

Ala Ala Ala Ser Ala Pro Val Tyr Gly Gln Ser Gly Leu Ala Tyr 65 70 75 80

Gly Pro Gly Ser Glu Ala Ala Ala Phe Gly Ala Asn Gly Leu Gly Gly 85 90 95

Phe Gln Pro Leu Asn Ser Val Ser Pro Ser Pro Leu Val Leu His 100 105 110

Pro Pro Pro Gln Leu Ser Pro Phe Leu His Pro His Gly Gln Gln Val 115 120 125

Pro Tyr Tyr Leu Glu Asn Glu Pro Ser Gly Tyr Ala Val Arg Glu Ala 130 135 140

Gly Pro Pro Ala Phe Tyr Arg Pro Asn Ser Asp Asn Arg Arg Gln Gly
145 150 155 160

25136771 - 50 -

Gly Arg Glu Arg Leu Ala Ser Thr Ser Asp Lys Gly Ser Met Ala Met 165 170 175

Glu Ser Ala Lys Glu Thr Arg Tyr Cys Ala Val Cys Asn Asp Tyr Ala 180 185 190

Ser Gly Tyr His Tyr Gly Val Trp Ser Cys Glu Gly Cys Lys Ala Phe 195 200 205

Phe Lys Arg Ser Ile Gln Gly His Asn Asp Tyr Met Cys Pro Ala Thr 210 215 220

Asn Gln Cys Thr Ile Asp Lys Asn Arg Arg Lys Ser Cys Gln Ala Cys 225 230 235 240

Arg Leu Arg Lys Cys Tyr Glu Val Gly Met Met Lys Gly Gly Ile Arg
245 250 255

Lys Asp Arg Arg Gly Gly Arg Met Leu Lys His Lys Arg Gln Arg Asp 260 265 270

Asp Gly Glu Gly Arg Asn Glu Ala Val Pro Pro Gly Asp Met Arg Ser 275 280 285

Ala Asn Leu Trp Pro Ser Pro Leu Leu Ile Lys His Thr Lys Lys Asn 290 295 300

Ser Pro Val Leu Ser Leu Thr Ala Asp Gln Met Ile Ser Ala Leu Leu 305 310 315 320

Glu Ala Glu Pro Pro Ile Ile Tyr Ser Glu Tyr Asp Pro Thr Arg Pro 325 330 335

Leu Ser Glu Ala Ser Met Met Gly Leu Leu Thr Asn Leu Ala Asp Arg 340 345 350

Glu Leu Val His Met Ile Asn Trp Ala Lys Arg Val Pro Gly Phe Leu 355 360 365

Asp Leu Ser Leu His Asp Gln Val His Leu Leu Glu Cys Ala Trp Leu 370 380

Glu Ile Leu Met Ile Gly Leu Val Trp Arg Ser Met Glu His Pro Gly 385 390 395 400

25136771 - 51 -

Lys Leu Leu Phe Ala Pro Asn Leu Leu Leu Asp Arg Asn Gln Gly Lys 405 410 415

Cys Val Glu Gly Met Val Glu Ile Phe Asp Met Leu Leu Ala Thr Ser 420 425 430

Ser Arg Phe Arg Met Met Asn Leu Gln Gly Glu Glu Phe Val Cys Leu 435 440 445

Lys Ser Ile Ile Leu Leu Asn Ser Gly Val Tyr Thr Phe Leu Ser Ser 450 460

Thr Leu Lys Ser Leu Glu Glu Lys Asp His Ile His Arg Val Leu Asp 465 470 475 480

Lys Ile Thr Asp Thr Leu Ile His Leu Met Ala Lys Ala Gly Leu Thr 485 490 495

Leu Gln Gln His Arg Arg Leu Ala Gln Leu Leu Leu Ile Leu Ser 500 505 510

His Phe Arg His Met Ser Asn Lys Gly Met Glu His Leu Tyr Asn Met 515 520 525

Lys Cys Lys Asn Val Val Pro Leu Tyr Asp Leu Leu Glu Met Leu 530 540

Asp Ala His Arg Leu His Ala Pro Thr Asn Leu Gly Gly Pro Pro 545 550 555 560

Glu Asp Met Ser Gln Ser Gln Leu Ala Thr Ser Gly Ser Thr Pro Ser 565 570 575

His Ser Leu Gln Met Tyr Tyr Ile Thr Gly Glu Ala Glu As<br/>n Phe Pro $580 \hspace{1.5cm} 585 \hspace{1.5cm} 590$ 

Thr Thr Ile 595

<210> 32

<211> 171

<212> PRT

<213> Human

<400> 32

Met Leu Leu Ser Ser Gln Val Asn Glu Glu His Ser Pro Val Ser Arg
1 5 10 15

25136771 - 52 -

Met Arg Thr Glu Phe Leu Met Gln Leu Asp Thr Val Leu Thr Ser Ala 20 Glu Asp Gln Ile Val Val Ile Cys Ala Thr Ser Lys Pro Glu Glu Ile Asp Glu Ser Leu Arg Arg Tyr Phe Met Lys Arg Leu Leu Ile Pro Leu Pro Asp Ser Thr Ala Arg His Gln Ile Ile Val Gln Leu Leu Ser Gln His Asn Tyr Cys Leu Asn Asp Lys Glu Phe Ala Leu Leu Val Gln Arg Thr Glu Gly Phe Ser Gly Leu Asp Val Ala His Leu Cys Gln Glu Ala 100 105 Val Val Gly Pro Leu His Ala Met Pro Ala Thr Asp Leu Ser Ala Ile 115 120 Met Pro Ser Gln Leu Arg Pro Val Thr Tyr Gln Asp Phe Glu Asn Ala 135 Phe Cys Lys Ile Gln Pro Ser Ile Ser Gln Lys Glu Leu Asp Met Tyr 150 155 Val Glu Trp Asn Lys Met Phe Gly Cys Ser Gln 165 <210> 33 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Estrogen Response Element <400> 33 gctctaagaa gaacagcctg

20

<210> 34 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Primer

<400> gctcta	34 agag gaacagcctg	20
<210><211><211><212><213>	DNA	
<220> <223>	Primer	
<400> agagag	35 atga tggggagggc aggggtgaag	30
<210><211><211><212><213>	36 6 DNA Artificial Sequence	
<220> <223>	Estrogen Response Element	
<400> aggtca	36	6
<210> <211> <212> <213>		
<220> <223>	Estrogen Response Element	
<400> tgacct	37	6
<210> <211> <212> <213>	38 13 DNA Artificial Sequence	
<220> <223>	Estrogen Response Element	
<222>	<pre>misc_feature (1)(13) n equals unknown</pre>	
<400> ggtcann	38 untg acc	13
<210> <211>	39 13	

25136771 - 54 -

<212> <213>	DNA Artificial Sequence	
<220> <223>	Estrogen Response Element	
<222>	<pre>misc_feature (1)(13) n equals unknown</pre>	
<400>	39	
aaccan	nntg act	13
<210><211><211><212><213>	5	
<220> <223>	Estrogen Response Element	
<400> ggtca	40	5
<210><211><211><212><213>	DNA	
<220> <223>	Estrogen Response Element	
<400> tggtc	41	5
<210> <211> <212> <213>	42 5 DNA Artificial Sequence	
<220> <223>	Estrogen Response Element	
<400> tgacc	42	5
<210><211><211><212><212><213>	43 22 DNA Artificial Sequence	
<220> <223>	Estrogen Response Element	
<400>	43	

25136771 - 55 -

attcga	tcag ggcggggcga gc	22
<210>	4.4	
<211>		
<212>		
	Artificial Sequence	
	•	
<220>		
<223>	Estrogen Response Element	
.000		
<220>		
<2212	misc_feature (1)(27)	
<223>	n equals unknown	
	1	
<400>	44	
gggcani	nnnn nnnnnnnnn nggcggg	27
<210>	45	
<211>	31	
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Estrogon Dognongo Element	
\2237	Estrogen Response Element	
<220>		
<221>	misc feature	
<222>	(1) $(31)$	
<223>	n equals unknown	
<400>	45	
		31
<b>J J</b>		31
<210>		
<211> <212>		
	Artificial Sequence	
\213/	Arctiticial Sequence	
<220>		
<223>	Estrogen Response Element	
<220>		
<221>	misc_feature (1)(23)	
	n equals unknown	
12237	ii equats unithown	
<400>	46	
gggccgc	gnn nnnnnnngg tca	23
<210>	47	
	5	
<212>	DNA	
<213>	Artificial Sequence	

```
<220>
<223> Estrogen Response Element
<400>
      47
gggca
                                                                      5
<210> 48
<211> 5
<212> DNA
<213> Artificial Sequence
<220>
<223>
      Estrogen Response Element
<400>
      48
ggtaa
                                                                      5
<210> 49
<211> 13
<212> DNA
<213> Artificial Sequence
<220>
<223> Estrogen Response Element
<220>
<221> misc_feature
<222> (1)..(13)
<223> n equals unknown
<400> 49
ggtcannntg ccc
                                                                    13
```

25136771 - 57 -